

Result No.	Score	Query Match	Length	DB	ID	Description
1	2072	100.0	2072	3	US-09-786-681A-1	Sequence 1, Appli
2	1814	87.5	1827	3	US-09-786-681A-3	Sequence 3, Appli
3	444	21.4	444	3	US-09-621-976-18829	Sequence 18829, A
4	383.4	18.5	440	3	US-09-513-993C-3708	Sequence 3708, A
5	369.8	17.8	771	3	US-09-270-767-679	Sequence 679, App
6	369.8	17.8	771	3	US-09-270-767-15961	Sequence 15961, A
7	364.8	17.6	433	3	US-09-513-993C-3502	Sequence 3502, A
8	230.8	11.1	571	3	US-09-270-767-28434	Sequence 28434, A
9	230.8	11.1	1151	3	US-09-270-767-12633	Sequence 12633, A
10	227.6	11.0	2391	3	US-09-949-016-3623	Sequence 3623, Ap
11	227.6	11.0	2805	3	US-08-959-004-6	Sequence 6, Appli
12	226	10.9	1878	3	US-10-104-047-1699	Sequence 1699, Ap
13	161.2	7.8	995	3	US-09-270-767-14715	Sequence 14715, A
14	132.8	6.4	726	3	US-09-248-796A-6208	Sequence 6208, Ap
15	122	5.9	499	3	US-09-949-016-1721	Sequence 1721, Ap
16	112.4	5.4	768	3	US-09-495-050A-10	Sequence 10, Appli
17	101	4.9	262	3	US-09-313-294A-2292	Sequence 2292, Ap
18	91.6	4.4	769	3	US-09-385-982-530	Sequence 530, App
19	79.8	3.9	227	3	US-09-490-609B-49	Sequence 49, Appl
20	70	3.4	601	3	US-09-949-016-59256	Sequence 59256, A
21	70	3.4	15148	3	US-09-949-016-13463	Sequence 13463, A
22	64.6	3.1	302	3	US-09-702-703-1002	Sequence 1002, Ap
23	64.6	3.1	302	3	US-09-736-457-1002	Sequence 1002, Ap
24	64.6	3.1	302	3	US-09-614-124B-1002	Sequence 1002, Ap

Db 301 AGTGGCTGGATATTAATAATTAAGATGATGATGCCAGCACTTACTGTGAATTTGAT 360  
Qy |||||  
Db 361 TTAGATAAGAAAAGAGAGATGCTATTTGTTATATGCCATAAAAAATCAATTAATGCTGACAG 420  
Qy |||||  
Db 361 TTAGATAAGAAAAGAGAGATGCTATTTGTTATATGCCATAAAAAATCAATTAATGCTGACAG 420  
Qy |||||  
Db 421 ATGTACATAGATGATTTACCAATATGGGGTATTTGTTGATGAGGCTGATGAAAATGGAGAA 480  
Qy |||||  
Db 421 ATGTACATAGATGATTTACCAATATGGGGTATTTGTTGATGAGGCTGATGAAAATGGAGAA 480  
Qy |||||  
Db 481 GATTACTATCTTTGACCTATAAAAACTTGAATAGTGTGTTTAAATGGAATCGAATTTGTT 540  
Qy |||||  
Db 481 GATTACTATCTTTGACCTATAAAAACTTGAATAGTGTGTTTAAATGGAATCGAATTTGTT 540  
Qy |||||  
Db 541 GATGTTAATCTAACTAGTGAAGGAAGGTGAACTGGTTCCAAATACTAAAATCCAGATG 600  
Qy |||||  
Db 541 GATGTTAATCTAACTAGTGAAGGAAGGTGAACTGGTTCCAAATACTAAAATCCAGATG 600  
Qy |||||  
Db 601 TCATATTCAGTAAATGGAATAAGTCAAGTGTGAAAATTTGAAGATCGAATTTGACAAATAT 660  
Qy |||||  
Db 601 TCATATTCAGTAAATGGAATAAGTCAAGTGTGAAAATTTGAAGATCGAATTTGACAAATAT 660  
Qy |||||  
Db 661 CTGTGATCCGTCCTTTTCAACATCGGATTCATTTGTTTCAATTTTCAACTCCCTTCATG 720  
Qy |||||  
Db 661 CTGTGATCCGTCCTTTTCAACATCGGATTCATTTGTTTCAATTTTCAACTCCCTTCATG 720  
Qy |||||  
Db 721 ATGTGATCTCTTGTGGGCTTAGTTTCAATGATTTTAAATGAGAACATTAAGAAAAGAT 780  
Qy |||||  
Db 721 ATGTGATCTCTTGTGGGCTTAGTTTCAATGATTTTAAATGAGAACATTAAGAAAAGAT 780  
Qy |||||  
Db 781 TATGCTCGGTACAGTAAAGGAAGAAATGATGATGATGATGATGATGATGATGATGATGATGAT 840  
Qy |||||  
Db 781 TATGCTCGGTACAGTAAAGGAAGAAATGATGATGATGATGATGATGATGATGATGATGATGAT 840  
Qy |||||  
Db 841 TATGATGGAAGAACAGGTGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 900  
Qy |||||  
Db 841 TATGATGGAAGAACAGGTGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 900  
Qy |||||  
Db 901 TCCTCTCTGATGTTCTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 960  
Qy |||||  
Db 901 TCCTCTCTGATGTTCTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 960  
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Qy |||||  
Db 961 GCAATGATAGATGATTTATATATCTGAGAGGGGATCAATGCTCAGTACAGCATATTTGTC 1020  
Qy |||||  
Db 1021 TATGCTCTGATCTCCAGTGAATGGTTTATTTGAGAGAGTCTGATGCTAGACAGGA 1080  
Qy |||||  
Db 1021 TATGCTCTGATCTCCAGTGAATGGTTTATTTGAGAGAGTCTGATGCTAGACAGGA 1080  
Qy |||||  
Db 1081 GGAAGGATGATGATTAAGACAGATGTTTATTTGGGCAATTCCTTATCCAGCTATGTTGTT 1140  
Qy |||||  
Db 1081 GGAAGGATGATGATTAAGACAGATGTTTATTTGGGCAATTCCTTATCCAGCTATGTTGTT 1140  
Qy |||||  
Db 1141 GGCATGCTCTTCTCATCAATTTCAATAGCCATTTATACCATGCTTCAAGGCCATTCCT 1200  
Qy |||||  
Db 1141 GGCATGCTCTTCTCATCAATTTCAATAGCCATTTATACCATGCTTCAAGGCCATTCCT 1200  
Qy |||||  
Db 1201 TTTGGAACAATGGTGGCGGTTGTTGATCTGTTTATTTGTTATTTCTCTCTTAAATCTT 1260  
Qy |||||  
Db 1201 TTTGGAACAATGGTGGCGGTTGTTGATCTGTTTATTTGTTATTTCTCTCTCTTAAATCTT 1260  
Qy |||||  
Db 1261 GTTGGTACAATCTATGCGGCAAAATCTGTGAGGTGAGCCCAACTTTCCTGTGCTGCAAT 1320  
Qy |||||  
Db 1261 GTTGGTACAATCTATGCGGCAAAATCTGTGAGGTGAGCCCAACTTTCCTGTGCTGCAAT 1320  
Qy |||||  
Db 1321 GCTGTGCTGCTCTATACCGAGAGAAAATGGTTTATGAGGCTGCGGTTATTTGTTG 1380  
Qy |||||  
Db 1321 GCTGTGCTGCTCTATACCGAGAGAAAATGGTTTATGAGGCTGCGGTTATTTGTTG 1380  
Qy |||||  
Db 1381 CTGGGTGGAATTTTACCTTTTGGTTCAATCTTTATGAAATGATTTTCACTTCCAGTCT 1440  
Qy |||||

Db 1381 CTGGGTGGAATTTTACCTTTTGGTTCAATCTTTATTTGAATGATTTTCACTTCCAGTCT 1440  
Qy |||||  
Db 1441 TTCTGGGCATATAAGATCTATTATGTTATGCTTCAATGATGCTGCTGCTGTTATCTCTG 1500  
Qy |||||  
Db 1441 TTCTGGGCATATAAGATCTATTATGTTATGCTTCAATGATGCTGCTGCTGTTATCTCTG 1500  
Qy |||||  
Db 1501 TGCATTTGACCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 1560  
Qy |||||  
Db 1501 TGCATTTGACCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 1560  
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Db 1561 TACCGGTGGCAATGGCAAGATTTTCTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 1620  
Qy |||||  
Db 1561 TACCGGTGGCAATGGCAAGATTTTCTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 1620  
Qy |||||  
Db 1621 TATTCCTTTTACTACTATTTTTCCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 1680  
Qy |||||  
Db 1621 TATTCCTTTTACTACTATTTTTCCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 1680  
Qy |||||  
Db 1681 TACTTTGGATATATGCGGTATTTAGCACAGCTTGGGATATGTTGAGGATGTTGTTGTTGTTG 1740  
Qy |||||  
Db 1681 TACTTTGGATATATGCGGTATTTAGCACAGCTTGGGATATGTTGAGGATGTTGTTGTTGTTG 1740  
Qy |||||  
Db 1741 TACATGGGAACAAGTCCCTTTTGTCCGAAAATCTATATACTGTAAGAAATTTGACTAGAGA 1800  
Qy |||||  
Db 1741 TACATGGGAACAAGTCCCTTTTGTCCGAAAATCTATATACTGTAAGAAATTTGACTAGAGA 1800  
Qy |||||  
Db 1801 CCCAGAAAACCTGGAACTTTTGGATCAATTTCTTTTCTAGGGGTGGAACCTTGCACAGC 1860  
Qy |||||  
Db 1801 CCCAGAAAACCTGGAACTTTTGGATCAATTTCTTTTCTAGGGGTGGAACCTTGCACAGC 1860  
Qy |||||  
Db 1861 AAAAAACAAAACCAAGAGAGATTTGGGCTTTAACTTTTTTTTTTTTTTTTTTTTTTTTTT 1920  
Qy |||||  
Db 1861 AAAAAACAAAACCAAGAGAGATTTGGGCTTTAACTTTTTTTTTTTTTTTTTTTTTTTTTT 1920  
Qy |||||  
Db 1921 TTTTTTTTTTTTTTTTACGAATGAGGCAATTTATTAACCCAGCATGTTGTTCTTAATGCT 1980  
Qy |||||  
Db 1921 TTTTTTTTTTTTTTTTACGAATGAGGCAATTTATTAACCCAGCATGTTGTTCTTAATGCT 1980  
Qy |||||  
Db 1981 TCTTGTGGCAGTGCACCTGTCCGGGATTTCTGTCAGATCTCTTTGCTCCCTGAGGTG 2040  
Qy |||||  
Db 1981 TCTTGTGGCAGTGCACCTGTCCGGGATTTCTGTCAGATCTCTTTGCTCCCTGAGGTG 2040  
Qy |||||  
Db 2041 TCAGTTTGGCGGCTCGAGCATGCACTAGA 2072  
Qy |||||  
Db 2041 TCAGTTTGGCGGCTCGAGCATGCACTAGA 2072  
Qy |||||

RESULT 2

US-09-786-681A-3  
; Sequence 3, Application US/09786681A  
; Patent No. 6692926  
; GENERAL INFORMATION:  
; APPLICANT: HIDAKA, Jun et al.  
; TITLE OF INVENTION: RECOMBINANT HUMAN SM-11044-BINDING RECEPTOR PROTEINS EXHIBITING L  
; TITLE OF INVENTION: BINDING ACTIVITIES, AND THEIR USES  
; FILE REFERENCE: 0020-4827P  
; CURRENT APPLICATION NUMBER: US/09/786,681A  
; CURRENT FILING DATE: 2001-01-24  
; NUMBER OF SEQ ID NOS: 7  
; SOFTWARE: PatentIn version 3.0  
; SEQ ID NO 3  
; LENGTH: 1827  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; FEATURE:  
; NAME/KEY: CDS  
; LOCATION: (11)..(1747)

Query Match 87.5%; Score 1814; DB 3; Length 1827;  
Best Local Similarity 100.0%; Pred. No. 0;  
Matches 1814; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Db	1094	GGCACTGCCCTTCCTTCATCAATTTTCATAGCCAAATTTATACCATGCTTCAAGAGCGAATTCCT	1153
Qy	1201	TTTGGAACAATGSGTGGCGGTTTGTCGATCTGTGTTTTTTTGTGTTATCTTCCTCTAAATCTTT	1260
Db	1154	TTTGGAACAATGSGTGGCGGTTTGTCGATCTGTGTTTTTTTGTGTTATCTTCCTCTAAATCTTT	1213
Qy	1261	GTTGGTACAATATCTTGGCCGAAATCTGTGAGGTCAGGTCAGCCAACTTTCTTCTGTGCTGTCAAT	1320
Db	1214	GTTGGTACAATCTTGGCCGAAATCTGTGAGGTCAGCCAACTTTCTTCTGTGCTGTCAAT	1273
Qy	1321	GCTGTGCTCGTCTCTATACCGGAGAAAAATGTTTCATGAGAGCTGGGGTTATTTGTTTGC	1380
Db	1274	GCTGTGCTCGTCTCTATACCGGAGAAAAATGTTTCATGAGAGCTGGGGTTATTTGTTTGC	1333
Qy	1381	CTGGGTGGAAATTTTACCTTTTGGTTCAACTCTTTATTTGAAATGTATTTTCATCTTCACGCTCT	1440
Db	1334	CTGGGTGGAAATTTTACCTTTTGGTTCAACTCTTTATTTGAAATGTATTTTCATCTTCACGCTCT	1393
Qy	1441	TTCTGGGCATATAGATCTATATATGTCATGCGCTTCATGATGCTGGTGCCTGCTTATCTCTG	1500
Db	1394	TTCTGGGCATATAGATCTATATATGTCATGCGCTTCATGATGCTGGTGCCTGCTTATCTCTG	1453
Qy	1501	TGCATTTGTGACTGTCTGTGTGACTATTTGTGTGCACATATTTTCTCTACTAAATGACAGAGAT	1560
Db	1454	TGCATTTGTGACTGTCTGTGTGACTATTTGTGTGCACATATTTTCTCTACTAAATGACAGAGAT	1513
Qy	1561	TACGGTGGCAATGGACAAGTTTTTCTCTCTGTGCAATCAACTGCAATCTATGTTTACATG	1620
Db	1514	TACGGTGGCAATGGACAAGTTTTTCTCTCTGTGCAATCAACTGCAATCTATGTTTACATG	1573
Qy	1621	TATTTCTTTTACTACTATTTTTCAAAAACAAGATGTATGGCTTATTTTCAAAATCATCATTT	1680
Db	1574	TATTTCTTTTACTACTATTTTTCAAAAACAAGATGTATGGCTTATTTTCAAAATCATCATTT	1633
Qy	1681	TACTTTGGATATATGGCGGTATTTAGCACAGCCTTGGGGATATATGTGTGGAGCGATGGGT	1740
Db	1634	TACTTTGGATATATGGCGGTATTTAGCACAGCCTTGGGGATATATGTGTGGAGCGATGGGT	1693
Qy	1741	TACATGGGAACAAGTGGCTTTGTCCGAAAAATCTATACTAATGTGAAAAATTGACTAGAGA	1800
Db	1694	TACATGGGAACAAGTGGCTTTGTCCGAAAAATCTATACTAATGTGAAAAATTGACTAGAGA	1753
Qy	1801	CCCAAGAAAAACCTGGAACTTTGGATCAATTTCTTTTTCATAGGGGTGGAACTTTGCACAGC	1860
Db	1754	CCCAAGAAAAACCTGGAACTTTGGATCAATTTCTTTTTCATAGGGGTGGAACTTTGCACAGC	1813
Qy	1861	AAAAACAACAAAC 1874	
Db	1814	AAAAACAACAAAC 1827	

RESULT 3  
US-09-621-976-18829  
: Sequence 18829. Application US/09621976

; Patent No. 6639063

**GENERAL INFORMATION:**

; APPLICANT: Dumas Milne Edwards, J.B.

**APPLICANT: Jobert, S.**

APPLICANT: Giordano, J.Y.

BYE REFERENCE: CENSET 250 ADDRESS AND NOTINENT: ESTB and

FILE REFERENCE: GENSET.034PK2  
CURRENT APPLICATION NUMBER: IIS/09/621 976

CURRENT AFFILIATION NUMBER: 05/0  
: CURRENT FILING DATE: 2000-07-21

NUMBER OF SEQ ID NOS: 19335

; SOFTWARE: Patent.pm

; SEQ ID NO 188

; LENGTH: 42

TYPE: DNA

ORGANISM: HOMO SAPIENS  
TTC-09-621-976-18829

63700T-0/E-T70-60-60

Query Match 21.4%; Score 444; DB 3; Length 444;

Best Local Similarity 100.0%; Pred. No. 1.5e-93;  
Matches 444; Conservative 0; Mismatches 0; Indels 0; Gaps 0;  
QY 550 CTAAGTGTGAGGAAAGGTAAGTGTGTTCCAAATTAATAATCCAGATGTCAATATCA 609  
Db 1 CTAAGTGTGAGGAAAGGTAAGTGTGTTCCAAATTAATAATCCAGATGTCAATATCA 60  
QY 610 GTAAATGGAAGGTAAGTGTGTTCCAAATTAATAATCCAGATGTCAATATCA 669  
Db 61 GTAAATGGAAGGTAAGTGTGTTCCAAATTAATAATCCAGATGTCAATATCA 120  
QY 670 TCCTTTTTCACATCGGATTCATGTTTTCCTCAATTTTCACTCCCTTCATGATGATC 729  
Db 121 TCCTTTTTCACATCGGATTCATGTTTTCCTCAATTTTCACTCCCTTCATGATGATC 180  
QY 730 TTTCTGGTGGCTTAGTTTCAATGATTTTAAATGAGAACATTAAGAAAGATTAATGCTCG 789  
Db 181 TTTCTGGTGGCTTAGTTTCAATGATTTTAAATGAGAACATTAAGAAAGATTAATGCTCG 240  
QY 790 TACAGTAAAGAGGAAAGGTAAGTGTGTTCCAAATTAATAATCCAGATGTCAATATCA 849  
Db 241 TACAGTAAAGAGGAAAGGTAAGTGTGTTCCAAATTAATAATCCAGATGTCAATATCA 300  
QY 850 AAACAGTGTGATGAGATGATTTAGACCATCAAGTCAACCCACTGATATTTTCTCTCTG 909  
Db 301 AAACAGTGTGATGAGATGATTTAGACCATCAAGTCAACCCACTGATATTTTCTCTCTG 360  
QY 910 ATTGGTTCGTGATGTCAGATATTTTGTCTGTCTCTCATCGTTATTTTGTGCAATGATA 969  
Db 361 ATTGGTTCGTGATGTCAGATATTTTGTCTGTCTCTCATCGTTATTTTGTGCAATGATA 420  
QY 970 GAAGATTTATATCTAGAGGGGA 993  
Db 421 GAAGATTTATATCTAGAGGGGA 444

RESULT 4

US-09-513-999C-3708  
Sequence 3708, Application US/09513999C  
Patent No. 6783961  
GENERAL INFORMATION:  
APPLICANT: Dumas Milne Edwards, J.B.  
APPLICANT: Duclert, A.  
APPLICANT: Giordano, J.Y.  
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.  
Patent No. 6783961  
FILE REFERENCE: 59.US2.REG  
CURRENT APPLICATION NUMBER: US/09/513,999C  
CURRENT FILING DATE: 2000-02-24  
PRIOR APPLICATION NUMBER: US 60/122,487  
PRIOR FILING DATE: 1999-02-26  
NUMBER OF SEQ ID NOS: 36681  
SOFTWARE: Patent.pm  
SEQ ID NO 3708  
LENGTH: 440  
TYPE: DNA  
ORGANISM: Homo sapiens  
FEATURE:  
NAME/KEY: CDS  
LOCATION: 180..440  
FEATURE:  
NAME/KEY: misc\_feature  
LOCATION: 151  
OTHER INFORMATION: m=a or c  
FEATURE:  
NAME/KEY: misc\_feature  
LOCATION: 155  
OTHER INFORMATION: s=g or c  
FEATURE:  
NAME/KEY: misc\_feature  
LOCATION: 162  
OTHER INFORMATION: k=g or t  
FEATURE:

NAME/KEY: misc\_feature  
LOCATION: 184  
OTHER INFORMATION: n=a, g, c or t  
FEATURE:  
NAME/KEY: misc\_feature  
LOCATION: 323  
OTHER INFORMATION: w=a or t  
FEATURE:  
NAME/KEY: misc\_feature  
LOCATION: 343  
OTHER INFORMATION: n=a, g, c or t  
FEATURE:  
NAME/KEY: misc\_feature  
LOCATION: 397  
OTHER INFORMATION: m=a or c  
FEATURE:  
NAME/KEY: misc\_feature  
LOCATION: 400  
OTHER INFORMATION: m=a or c  
FEATURE:  
NAME/KEY: UNSURE  
LOCATION: 2  
OTHER INFORMATION: Xaa=Lys or Met or Arg or Thr  
FEATURE:  
NAME/KEY: UNSURE  
LOCATION: 55  
OTHER INFORMATION: Xaa=Ala or Asp or Gly or Val  
FEATURE:  
NAME/KEY: UNSURE  
LOCATION: 73  
OTHER INFORMATION: Xaa=Ala or Asp  
FEATURE:  
NAME/KEY: UNSURE  
LOCATION: 74  
OTHER INFORMATION: Xaa=Lys or Thr  
US-09-513-999C-3708

Query Match 18.5%; Score 383.4; DB 3; Length 440;  
Best Local Similarity 95.7%; Pred. No. 1.8e-79;  
Matches 420; Conservative 5; Mismatches 8; Indels 6; Gaps 3;  
QY 638 TTGAAGATCGATTTGACAAATATCTTGATCCGCTCTTTTTCACACATCGGATTCATG 697  
Db 2 TTGAAGATCGATTTGACAAATATCTTGATCCGCTCTTTTTCACACATCGGATTCATG 61  
QY 698 TTTCAATTTTCACTCTTCATGATGATGATCTTCTTGTTGGCTTAGTTTCAATGATTT 757  
Db 62 TTTCAATTTTCACTCTTCATGATGATGATCTTCTTGTTGGCTTAGTTTCAATGATTT 121  
QY 758 TAATGAGAACATTAAGAAAG---ATTATGCTCGTACAGTAAAGAGGAAAGATGGAT 813  
Db 122 TAATGAGAACATTAAGAAAGAAATTAATGCTCGTACAGTAAAGAGGAAAGATGGAT 181  
QY 814 GAT-ATGATAGACACCTAGGAGATGAATATGATGGAACAGGTGTCATGAGATGATTT 872  
Db 182 GAGATGATAGACACCTAGGAGATGAATATGATGGAACAGGTGTCATGAGATGATTT 241  
QY 873 TAGACCATCAAGTCAACCACTGATATTTTCTCTCTGATGTTGTTCTGATGTCAGATATTT 932  
Db 242 TAGACCATCAAGTCAACCACTGATATTTTCTCTCTGATGTTGTTCTGATGTCAGATATTT 301  
QY 933 TGCTGTGCTCTCATCGTTATTTATTTGTCATGATGATGATGATTTATATCTAGAGGGG 992  
Db 302 TGCTGTGCTCTCATCGTTATTTATTTGTCATGATGATGATGATTTATATCTAGAGGGG 361  
QY 993 ATCAATGCTCAGTACAGGCAATATTTGTTCTATGCTCTGATGCT-CCAGTGAATGGTATTT 1051  
Db 362 ATCAATGCTCAGTACAGGCAATATTTGTTCTATGCTCTGATGCTCTCCAGTGAATGGTATTT 421  
QY 1052 TTGGAGGAAGTCTGTATGC 1070  
Db 422 TTGGAGGAAGTCTGTATGC 440



RESULT 5  
US-09-270-767-679/c  
; Sequence 679, Application US/09270767  
; Patent No. 6703491  
; GENERAL INFORMATION:  
; APPLICANT: Homburger et al.  
; TITLE OF INVENTION: Nucleic acids and proteins of *Drosophila melanogaster*  
; FILE REFERENCE: File Reference: 7326-094  
; CURRENT APPLICATION NUMBER: US/09/270,767  
; CURRENT FILING DATE: 1999-03-17  
; NUMBER OF SEQ ID NOS: 62517  
; SOFTWARE: PatentIn Ver. 2.0  
; SEQ ID NO 679  
; LENGTH: 771  
; TYPE: DNA  
; ORGANISM: *Drosophila melanogaster*  
US-09-270-767-679

Query Match 17.8%; Score 369.8; DB 3; Length 771;  
Best Local Similarity 67.7%; Pred. No. 3.2e-76;  
Matches 518; Conservative 0; Mismatches 247; Indels 0; Gaps 0;

QY 610 GTAAATGGAAGAGTCAAGATGCAAAATTTGAAGATCGATTTGACAAATATCTTGATCCG 669  
DB 765 GTCNACTGGAGCCAGCAGAGTGGAGTTCAAGATCGATTCGACAAAGTACCTGGATCCC 706  
QY 670 TCCTTTTTCACACATCGGATTCATTTGTTTTCATTTTCAACTCTTTCATGATGGTGATC 729  
DB 705 AACTTCTTCAGACACAGATCCACTGGTTCAGCATCTTCAACAGCTTTCATGATGGTGATC 646  
QY 730 TTCTTGTGGCTTAGTTTCAATGATTTTAAATGAGAACATTAAGAAAGATTTATGCTCGG 789  
DB 645 TTCTTGTGGCTTAGTTTCAATGATTTTAAATGAGAACATTAAGAAAGATTTATGCTCGG 586  
QY 790 TACAGTAAAGAGAGAGAAATGATGATGATGATGATGATGATGATGATGATGATGATG 849  
DB 585 TACAGTAAAGAGAGAGAAATGATGATGATGATGATGATGATGATGATGATGATGATG 526  
QY 850 AAGCAGGTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATG 909  
DB 525 AAGCAGGTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATG 466  
QY 910 ATTGGTCTCGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATG 969  
DB 465 GTGGGCGCTCGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATG 406  
QY 970 GAAGATTTATATCTAGAGAGGAGATCAATGCTCAGTACAGCATATTTGCTATGCTGCT 1029  
DB 405 GGTGAATTTGATACAGGAGCGGCTCAATGCTGCTCAGGCTATATTTGATGATGCGCC 346  
QY 1030 ACGTCTCCAGTCAATGTTTATTTGAGGAGAGTCTGTATGCTAGACAGAGAGAGAGAGA 1089  
DB 345 ACCTCACCATCAATGATGATGATGATGATGATGATGATGATGATGATGATGATGATG 286  
QY 1090 TGGATCCGACAGATGCTGGTGCTGCTTTTACAGTTCCAGTGGCTGTGTGCGGACCGCT 1149  
DB 285 TGGATCCGACAGATGCTGGTGCTGCTTTTACAGTTCCAGTGGCTGTGTGCGGACCGCT 226  
QY 1150 TTCTTCATCAATTTATAGCATTATTTACATGCTTCAAGAGCCATTCCTTTTGAACA 1209  
DB 225 TTCTTCATCAATTTATAGCATTATTTACATGCTTCAAGAGCCATTCCTTTTGAACA 166  
QY 1210 ATGGTGCGCGTTTGTGATCTGTTTTTGTATTTTCTTCTAAATCTGTTGTGTGATA 1269  
DB 165 ATGGTGCGCGTTTGTGATCTGTTTTTGTATTTTCTTCTAAATCTGTTGTGTGATA 106  
QY 1270 ATACTTGGCCGAATCTGTGATGATGATGATGATGATGATGATGATGATGATGATG 1329  
DB 105 GTCTGCGCGCGAATCTGTGATGATGATGATGATGATGATGATGATGATGATGATG 1374  
QY 1330 CGTCTTATACCGAGAGAAATGTTTATGATGATGATGATGATGATGATGATGATGATG 1374

Db 45 CGACCCATTCCCGAAAAAGAGTGGTATCATGAGCCACTGATTATT 1

RESULT 6  
US-09-270-767-15961/c  
; Sequence 15961, Application US/09270767  
; Patent No. 6703491  
; GENERAL INFORMATION:  
; APPLICANT: Homburger et al.  
; TITLE OF INVENTION: Nucleic acids and proteins of *Drosophila melanogaster*  
; FILE REFERENCE: File Reference: 7326-094  
; CURRENT APPLICATION NUMBER: US/09/270,767  
; CURRENT FILING DATE: 1999-03-17  
; NUMBER OF SEQ ID NOS: 62517  
; SOFTWARE: PatentIn Ver. 2.0  
; SEQ ID NO 15961  
; LENGTH: 771  
; TYPE: DNA  
; ORGANISM: *Drosophila melanogaster*  
US-09-270-767-15961

Query Match 17.8%; Score 369.8; DB 3; Length 771;  
Best Local Similarity 67.7%; Pred. No. 3.2e-76;  
Matches 518; Conservative 0; Mismatches 247; Indels 0; Gaps 0;

QY 610 GTAAATGGAAGAGTCAAGATGCAAAATTTGAAGATCGATTTGACAAATATCTTGATCCG 669  
DB 765 GTCNACTGGAGCCAGCAGAGTGGAGTTCAAGATCGATTCGACAAAGTACCTGGATCCC 706  
QY 670 TCCTTTTTCACACATCGGATTCATTTGTTTTCATTTTCAACTCTTTCATGATGGTGATC 729  
DB 705 AACTTCTTCAGACACAGATCCACTGGTTCAGCATCTTCAACAGCTTTCATGATGGTGATC 646  
QY 730 TTCTTGTGGCTTAGTTTCAATGATTTTAAATGAGAACATTAAGAAAGATTTATGCTCGG 789  
DB 645 TTCTTGTGGCTTAGTTTCAATGATTTTAAATGAGAACATTAAGAAAGATTTATGCTCGG 586  
QY 790 TACAGTAAAGAGAGAGAAATGATGATGATGATGATGATGATGATGATGATGATGATG 849  
DB 585 TACAGTAAAGAGAGAGAAATGATGATGATGATGATGATGATGATGATGATGATGATG 526  
QY 850 AAGCAGGTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATG 909  
DB 525 AAGCAGGTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATG 466  
QY 910 ATTGGTCTCGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATG 969  
DB 465 GTGGGCGCTCGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATG 406  
QY 970 GAAGATTTATATCTAGAGAGGAGATCAATGCTCAGTACAGCATATTTGCTATGCTGCT 1029  
DB 405 GGTGAATTTGATACAGGAGCGGCTCAATGCTGCTCAGGCTATATTTGATGATGCGCC 346  
QY 1030 ACGTCTCCAGTCAATGTTTATTTGAGGAGAGTCTGTATGCTAGACAGAGAGAGAGAGA 1089  
DB 345 ACCTCACCATCAATGATGATGATGATGATGATGATGATGATGATGATGATGATGATG 286  
QY 1090 TGGATCCGACAGATGCTGGTGCTGCTTTTACAGTTCCAGTGGCTGTGTGCGGACCGCT 1149  
DB 285 TGGATCCGACAGATGCTGGTGCTGCTTTTACAGTTCCAGTGGCTGTGTGCGGACCGCT 226  
QY 1150 TTCTTCATCAATTTATAGCATTATTTACATGCTTCAAGAGCCATTCCTTTTGAACA 1209  
DB 225 TTCTTCATCAATTTATAGCATTATTTACATGCTTCAAGAGCCATTCCTTTTGAACA 166  
QY 1210 ATGGTGCGCGTTTGTGATCTGTTTTTGTATTTTCTTCTAAATCTGTTGTGTGATA 1269  
DB 165 ATGGTGCGCGTTTGTGATCTGTTTTTGTATTTTCTTCTAAATCTGTTGTGTGATA 106  
QY 1270 ATACTTGGCCGAATCTGTGATGATGATGATGATGATGATGATGATGATGATGATG 1329  
DB 105 GTCTGCGCGCGAATCTGTGATGATGATGATGATGATGATGATGATGATGATGATG 46

QY 1330 CGTCTATACCGAGAAAAATGTTTCATGAGCGCTCGGTTATT 1374  
Db 45 CGACCCATTCCGNAAGAGATGGTATACATGAGCGCACTGATTATT 1

## RESULT 7

US-09-513-999C-3502  
; Sequence 3502, Application US/09513999C  
; Patent No. 6783961  
; GENERAL INFORMATION:  
; APPLICANT: Dumas Milne Edwards, J.B.  
; APPLICANT: Duclert, A.  
; APPLICANT: Giordano, J.Y.  
; TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.  
; Patent No. 6783961  
; FILE REFERENCE: 59.US2.REG  
; CURRENT APPLICATION NUMBER: US/09/513,999C  
; CURRENT FILING DATE: 2000-02-24  
; PRIOR APPLICATION NUMBER: US 60/122,487  
; PRIOR FILING DATE: 1999-02-26  
; NUMBER OF SEQ ID NOS: 36681  
; SOFTWARE: Patent.pm  
; SEQ ID NO 3502  
; LENGTH: 433  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; FEATURE:  
; NAME/KEY: CDS  
; LOCATION: 100..432  
; FEATURE:  
; NAME/KEY: misc\_feature  
; LOCATION: 86  
; OTHER INFORMATION: m=a or c  
US-09-513-999C-3502

Query Match 17.6%; Score 364.8; DB 3; Length 433;  
Best Local Similarity 98.9%; Pred. No. 3.8e-75;  
Matches 366; Conservative 1; Mismatches 3; Indels 0; Gaps 0;  
QY 124 CACAGGTATCAAGATAAAGAGGAGTGTCTTATGATGAATACTGTGGGCGCTTACCAT 183  
Db 64 CAGCAGTATCAAGATAAAGAGGAGTGTCTTATGATGAATACTGTGGGCGCTTACCAT 123  
QY 184 AATCGTCAAGACATATAGTACTTTTCTACTTCCATCTCTGTGGGTCAGAAAAAAGT 243  
Db 124 AATCGTCAAGACATATAGTACTTTTCTACTTCCATCTCTGTGGGTCAGAAAAAAGT 183  
QY 244 ATCAGTCATTACCATGAAACTCTGGGAGAGCACTTCAAGGGGTGAAATTTAGT 303  
Db 184 ATCAGTCATTACCATGAAACTCTGGGAGAGCACTTCAAGGGGTGAAATTTAGT 243  
QY 304 GGTCTGATATTAATTTAAAGATGATGTGATGCCAGCCACTTACTGTGAAATGATTTA 363  
Db 244 GGTCTGATATTAATTTAAAGATGATGTGATGCCAGCCACTTACTGTGAAATGATTTA 303  
QY 364 GATAGAGAGAGAGATGATTTGTTATATGCCATAAATAATCATCTACTGTACCAAGATG 423  
Db 304 GATAGAGAGAGAGATGATTTGTTATATGCCATAAATAATCATCTACTGTACCAAGATG 363  
QY 424 TACATAGATGATTTACCAATATGGGTATTTGTTGGTGAGGCTGATGAAATGGAGAAGAT 483  
Db 364 TACATAGATGATTTACCAATATGGGTATTTGTTGGTGAGGCTGATGAAATGGAGAAGAT 423  
QY 484 TACTATCTTT 493  
Db 424 TACTATCTTT 433

## RESULT 8

US-09-270-767-28434  
; Sequence 28434, Application US/09270767  
; Patent No. 6703491  
; GENERAL INFORMATION:

; APPLICANT: Homburger et al.  
; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster  
; FILE REFERENCE: File Reference: 7326-094  
; CURRENT APPLICATION NUMBER: US/09/270,767  
; CURRENT FILING DATE: 1999-03-17  
; NUMBER OF SEQ ID NOS: 62517  
; SOFTWARE: PatentIn Ver. 2.0  
; SEQ ID NO 28434  
; LENGTH: 571  
; TYPE: DNA  
; ORGANISM: Drosophila melanogaster  
US-09-270-767-28434

Query Match 11.1%; Score 230.8; DB 3; Length 571;  
Best Local Similarity 72.7%; Pred. No. 5.9e-44;  
Matches 298; Conservative 0; Mismatches 112; Indels 0; Gaps 0;  
QY 1394 TACCTTTTGGTTCAATCTTTTATTGAAATGATTTTCACTTTCAGCTTTTCTGGGCATATA 1453  
Db 1 TGCCCTTTGGATCCATCTTCAATTGAGATGATCTTCACTTTCACCTCTTCTGGGCATATA 60  
QY 1454 AGATCTATTATGCTCTATGGCTTCATGATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 1513  
Db 61 AGATCTACTACGCTTACGGCTTCTATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 120  
QY 1514 TCTGTGTGACTATTGTGTGCACATATTTTCTACTAAATGCAAGAGATTACCGGTGGCAAT 1573  
Db 121 TGTGGTCAACATGCTGTGTGCACCTTCTCTCTTAATGCCAGGATTAACGATGGCAGT 180  
QY 1574 GGACAAAGTTTCTCTCTGCTGCATCAACTGCAATCTATGTTTACATGATTTCTTTTACT 1633  
Db 181 GGACGAGTTTTCATGGCTGGGGCTCCACGTCGATTTACGTAGCGCTATTCTCTCTATT 240  
QY 1634 ACTATTTTCAAAACAAGATGATGGCTTATTTTCAACATCATTTTACTTTTGGATATA 1693  
Db 241 ACTTCTTCTTTAAACCAAAATGTTGGGCTGTTCCTCAACGGCTTCTTACTTTTGGCTACA 300  
QY 1694 TGCGCGTATTTAGACAGCCTTTGGGGATTAATGTGTGGAGCGATTGGTTACATGGGAACAA 1753  
Db 301 TGCGACTCTTTCAGCGCGCTTTGGGCATTTATCTGCGGCACCGCTGGCTATGTGGGCACGA 360  
QY 1754 GTGCCCTTTCGCAAAATCTATATCTAATGTGAAATTTGCAATGACATAGAGACCC 1803  
Db 361 ATCTCTTTGGCGCAAAATCTATTCCAATGTGAAATAGACTAAGAGCCC 410

## RESULT 9

US-09-270-767-12633  
; Sequence 12633, Application US/09270767  
; Patent No. 6703491  
; GENERAL INFORMATION:  
; APPLICANT: Homburger et al.  
; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster  
; FILE REFERENCE: File Reference: 7326-094  
; CURRENT APPLICATION NUMBER: US/09/270,767  
; CURRENT FILING DATE: 1999-03-17  
; NUMBER OF SEQ ID NOS: 62517  
; SOFTWARE: PatentIn Ver. 2.0  
; SEQ ID NO 12633  
; LENGTH: 1151  
; TYPE: DNA  
; ORGANISM: Drosophila melanogaster  
US-09-270-767-12633

Query Match 11.1%; Score 230.8; DB 3; Length 1151;  
Best Local Similarity 72.7%; Pred. No. 7.6e-44;  
Matches 298; Conservative 0; Mismatches 112; Indels 0; Gaps 0;  
QY 1394 TACCTTTTGGTTCAATCTTTTATTGAAATGATTTTCACTTTCAGCTTTTCTGGGCATATA 1453  
Db 1 TGCCCTTTGGATCCATCTTCAATTGAGATGATCTTCACTTTCACCTCTTCTGGGCATATA 60  
QY 1454 AGATCTATTATGCTCTATGGCTTTCATGATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 1513

Db 61 AGATCTACTAGCTTACGGCTTCATGTTGCTGGTTCAGCATCCTGACTGTCACCG 120  
Qy 1514 TCTGTGACATATTGTGTGACATATTTTCTACTAAATGACAGATTTACCGTGGCAAT 1573  
Db 121 TGTGGTCAACATCGTGTGACCTACTTCTGCTAAATGCCGAGATTAACGATGGCAGT 180  
Qy 1574 GGACAAGTTTCTCTGCTGCATCACTGCAATCTATGTTTATACATGATTTCTTTTACT 1633  
Db 181 GGACGAGTTTCATGCTCGGGCTCCACGTCGATTAAGTGTAGTGGCTTATTCCTTCTATT 240  
Qy 1634 ACTATTTTTCACAAACAAAGATGATGCTTATTTTCAACATCATTTTACTTTGATATA 1693  
Db 241 ACTTCTTTTAAACCAAAATGTTGGTCTGTGTTCCAAACGGCTTCTACTTTGGCTACA 300  
Qy 1694 TGGCGGTATTTAGCACAGCTTGGGGAATAATGTGTGGAGCAATTTGTTACATGGGAACAA 1753  
Db 301 TGGCACTCTTCAGCGGCGCTTGGGCATTAATCTGGGCGACCGTCTGCTATGTGGGCAAG 360  
Qy 1754 GTGCTTTTGTCCGAAATCTATCTAATGTGAAATTTGACTAGAGACC 1803  
Db 361 ATCTCTTTGTGCGCAAAATCTATTCCTCAATGTGAAATAGACTAAGAGCCC 410

## RESULT 10

US-09-949-016-3623  
; Sequence 3623, Application US/09949016  
; Patent No. 6812339  
; GENERAL INFORMATION:  
; APPLICANT: VENTER, J. Craig et al.  
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED  
; FILE REFERENCE: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF  
; CURRENT APPLICATION NUMBER: US/09/949,016  
; PRIOR FILING DATE: 2000-04-14  
; PRIOR APPLICATION NUMBER: 60/241,755  
; PRIOR FILING DATE: 2000-10-20  
; PRIOR APPLICATION NUMBER: 60/237,768  
; PRIOR FILING DATE: 2000-10-03  
; PRIOR APPLICATION NUMBER: 60/231,498  
; PRIOR FILING DATE: 2000-09-08  
; NUMBER OF SEQ ID NOS: 207012  
; SOFTWARE: FastSeq for Windows Version 4.0  
; SEQ ID NO 3623  
; LENGTH: 2391  
; TYPE: DNA  
; ORGANISM: Human  
US-09-949-016-3623

Query Match 11.0%; Score 227.6; DB 3; Length 2391;  
Best Local Similarity 51.9%; Pred. No. 5.5e-43;  
Matches 596; Conservative 0; Mismatches 534; Indels 18; Gaps 3;  
Qy 656 AATATCTTGATCCGCTCTTTTCAACATCGATTCATTTGTTTCAATTTTCACTTCT 715  
Db 996 ACTATATTTCTGGAGTCTATGCTCTATCCCATCCCATCTAGTGGTTAGCATTAATGAATTC 1055  
Qy 716 TCATGATGTTGATCTTCTTCTGCTGGCTTATGTTTCAATGATTTTAAATGAGAACATTAAGAA 775  
Db 1056 TGGTCATGTTCTTCTTCTATCTGATGGTATGATGATGATGATGATGATGATGATGATGATG 1115  
Qy 776 AAGATTATGCTCGGTACAGTAAAGAGGAGAAATGGATGATGATGATGATGATGATGATGATGAT 835  
Db 1116 AAGATATTTGCTAGATATAATCAGATGGACTCTACGGAAGATGCCAG-----G 1163  
Qy 836 ATGATATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 895  
Db 1164 AAGAAATTTGGCTGGAATCTTCTCATGTTGATATATTCGCTCCCAAGAAAGGGATGC 1223  
Qy 896 TATTTTCTCTCTGATGTTGTTCTGATGTCAGATGATGATGATGATGATGATGATGATGATGAT 955  
Db 1224 TGCTATGATCTTTCTAGGATCCGGGACACAGATTTTAAATGATGATGATGATGATGATGATGAT 1283

Qy 956 TTCTTGCAATGATGAGAAATTTATATATCTGAGAGGGATCATATGCTCAGTAC---AGCCA 1012  
Db 1284 TTTTTCGCTTGGCTGGGATTTTGTCTACCTGCCAACGAGAGGCTGATGACGTGCTG 1343  
Qy 1013 TATTTGCTATGCTGCTCCTCCTCAGTGAATGTTTATTTTGGAGGAAGTCTGTATGCTA 1072  
Db 1344 TGTCTCTGTTGGTGTCTGCTGGGCACTCCCTGCGAGCTATGTTGCTGCCAGATTTCTTAAGT 1403  
Qy 1073 GACAGAGGAGGAGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1132  
Db 1404 CTTTGGAGGTGAGAAAGTGGAAACAAATGTTTATTAACATCAATTTCTTTGCTCTGGGA 1463  
Qy 1133 TGTGTTGTCACATGCTCTTCTTCTCATCAATTTATAGCCATTTTATTAACATGCTTCAAGAG 1192  
Db 1464 TTGTAATTTGCTGACTTCTTTTAAATGAATCTGATCTCTGCGGAGAGAGATCTTCAGCAG 1523  
Qy 1193 CCATTCCTTTTGGAAACAATGTTGGCGGTTTGTGTCATCTGTTTGTGTTTGTGTTTGTGTTT 1252  
Db 1524 CTATTCCTTTTGGGACATGTTGTCATATGTCCTCTTGGCTCTGTCATATCTGTGCTC 1583  
Qy 1253 TAAATCTGTTGTTGATCAATCTGTTGGCGGAAATCTGTGAGTCAAGCCCACTTTCTTGTGTC 1312  
Db 1584 TGACGTTTATTTGCTGATCTTTTGGTGTGTTTAAAGAAAGATGCCATTTGAACAC---CCAGTTC 1640  
Qy 1313 GTGTCATGCTGCTGCTGCTCTCTATACCGGAGAAATGTTTCTGTCGAGCCCTGCGGTTA 1372  
Db 1641 GAACCAATCAGATTTCCACGCTCAGATTTCTGAAACAGTCTGTTCTACAGAAACCTTGGCTG 1700  
Qy 1373 TTGTTTGGCTTGGTGGAAATTTTACCTTTTGGTTTCAATCTTTTATTTGAAATGATTTTCACT 1432  
Db 1701 GTATTAATCATGAGGAGGATTTTGGCTTGGCTGTCATCTTTATACAACTTTTCTTCACTC 1760  
Qy 1433 TCACGCTTTTCTGGGCAATTAAGATCTATATGCTATGCTTTCATGATGCTGCTGCTGCTG 1492  
Db 1761 TGAATAGTATTTGGTCAACACAGATGATTTACATGTTTGGCTTCTTATTTCTGCTGTTA 1820  
Qy 1493 TTATCTGTCATTTGTCATGCTGCTGTCATGTCATGTCATGTCATGTCATGTCATGTCATGTC 1552  
Db 1821 TCATTTTGGTATTTACCTGTTCTGAAAGCAACTATCTCTTCTGCTATTTTCCACCTATGTC 1880  
Qy 1553 CAGAAGATTTACCGTGGCAATGGAACAAGTTTCTCTGCTGTCATCAATGCAATCTATG 1612  
Db 1881 CAGAGATTTATCATTTGGCAATGCGTTCATCTCTACGAGTGGCTTTTACTGCGATTTATT 1940  
Qy 1613 TTTACATGATTTCTTTTACTACTATTTTTCACAAACAAGATGATGCTGCTGCTGCTGCTGCT 1672  
Db 1941 TCTTAATCTATGCTAGTACACTACTTCTTTTCAAAACCTGACAGATCAGCGGAAACAGCA 2000  
Qy 1673 CATCATTTTACTTTTGGATATATGCGGTATTTAGCACAGCCTTGGGATTAATGTCGAG 1732  
Db 2001 CAATTCGTACTTTGTTTATACCATGATTAATGTTTGTGATCTTCTTCTTTTACAGGAA 2060  
Qy 1733 CGATTCGTTACATGGAACAAGTGGCTTTTGTCCGAAAAATCTATACATAATGTCGAAAAATG 1792  
Db 2061 CAATTCGCTTCTTGGATGCTTTTGGTGTGTTTGTGTACCAAAATATACAGTGTGTGAGGTTG 2120  
Qy 1793 ACTAGAGA 1800  
Db 2121 ACTGAAGA 2128

## RESULT 11

US-08-959-004-6  
; Sequence 6, Application US/08959004  
; Patent No. 6197543  
; GENERAL INFORMATION:  
; APPLICANT: Hillman, Jennifer L.  
; APPLICANT: Yue, Henry  
; APPLICANT: Corley, Neil C.  
; APPLICANT: Lal, Preeti  
; APPLICANT: Shah, Purvi  
; APPLICANT: Kaser, Matthew  
; TITLE OF INVENTION: HUMAN VESICLE MEMBRANE PROTEIN-LIKE

;; TITLE OF INVENTION: PROTEINS  
;; NUMBER OF SEQUENCES: 11  
;; CORRESPONDENCE ADDRESS:  
;; ADDRESS: Incyte Pharmaceuticals, Inc.  
;; STREET: 3174 Porter Drive  
;; CITY: Palo Alto  
;; STATE: CA  
;; COUNTRY: USA  
;; ZIP: 94304  
;; COMPUTER READABLE FORM:  
;; MEDIUM TYPE: Diskette  
;; COMPUTER: IBM Compatible  
;; OPERATING SYSTEM: DOS  
;; SOFTWARE: FASSEQ for Windows Version 2.0  
;; CURRENT APPLICATION DATA:  
;; APPLICATION NUMBER: US/08/959,004  
;; FILING DATE: Herewith  
;; CLASSIFICATION: 514  
;; PRIOR APPLICATION DATA:  
;; APPLICATION NUMBER:  
;; FILING DATE:  
;; ATTORNEY/AGENT INFORMATION:  
;; NAME: Billings, Lucy J.  
;; REGISTRATION NUMBER: 36,749  
;; REFERENCE/DOCKET NUMBER: PF-0414 US  
;; TELECOMMUNICATION INFORMATION:  
;; TELEPHONE: 650-855-0555  
;; TELEFAX: 650-845-4166  
;; TELEX:  
;; INFORMATION FOR SEQ ID NO: 6:  
;; SEQUENCE CHARACTERISTICS:  
;; LENGTH: 2805 base pairs  
;; TYPE: nucleic acid  
;; STRANDEDNESS: single  
;; TOPOLOGY: linear  
;; IMMEDIATE SOURCE:  
;; LIBRARY: ADREUT06  
;; CLONE: 2822412  
;; US-08-959-004-6

Query Match 11.0%; Score 227.6; DB 3; Length 2805;  
Best Local Similarity 51.9%; Pred. No. 5.8e-43;  
Matches 596; Conservative 0; Mismatches 534; Indels 18; Gaps 3;

Qy 656 AATATCTTGATCCGCTCTTTTCAACATCGGATCATGTGTTTCAATTTCAACTCCT 715  
Db 1044 ACTATATCTGGAGTCTATGCTCATACCACATTCAGTCTTAGCATTAATGAATCCC 1103  
Qy 716 TCATGATGCTGATCTTCTTGGTGGCTTAGTTTCAATGATTTTAATGAGACATTAAGAA 775  
Db 1104 TGGTCAATGTTCTCTTCTTATCTGGAATGGTAGCTATGATTTATGTTACGGACCTGCACA 1163  
Qy 776 AAGATTATGCTCGGTACAGTAAAGAGGAAGAAATGGATGATGATGATGAGACCTAGGAG 835  
Db 1164 AAGATATTGCTAGATATAATCAGATGGACTCTACGGAAGATGCCAG-----G 1211  
Qy 836 ATGAATATGATGGAACAGAGTGCATGAGATGATTTATGACCATCAATCAAGTCAACCATGA 895  
Db 1212 AAGAAATTTGGCTGGAACTTTGTCATGTGTATATATTCGCTCCCAAGAAAGGGATGC 1271  
Qy 896 TATTTCTCTCTGATCTGATGCTGATGCTGATGCTGATGCTGCTCTCATCGTTATTA 955  
Db 1272 TGCTATCAGTCTTCTAGATCCGGGACACAGATTTTAATATGACCTTTGTGACTAT 1331  
Qy 956 TTGTTGCAATGATGAAGATTTATATCTGAGAGGGGATCAATGCTCAGTAC---AGCCA 1012  
Db 1332 TTTTGGCTTGGGATTTTGTACCTGCCACCGAGGAGCGCTGATGAGGTGCTG 1391  
Qy 1013 TATTTGCTATGCTGCTACGCTCCAGTGAATGTTATTTGGAGGAAGTCTGTATGCTA 1072  
Db 1392 TGGTCTGTGGGTGCTGCTGGGCACCCCTGCAGGCTATGTTGCTGCCAGATTTCTATAAGT 1451  
Qy 1073 GACAAGGAGGAGGAGATGAGTAAAGCAGATGTTTATTTGGGGCATTCCTTATCCAGCTA 1132

Db 1452 CCTTGGAGGTGAGAGTGGAAAACAAATGTTTATTAACATCATTTCTTTGTCTCTGGGA 1511  
Qy 1133 TGGTGTGTGCACTGCCCTTCTTCATCAATTTTCATAGCCATTTATTAACATGCTTTCAAGAG 1192  
Db 1512 TTGTAATTTGCTGACTTCTTTTATAATGAATCTGATCCTCTCGGGGAGAGGATCTTTCAGCAG 1571  
Qy 1193 CCAATTCCTTTTGGAAACATGSGTGGCGGTTTGTGTGCATCTGTTTGTGTTTATTTCTTCTC 1252  
Db 1572 CTATTCCTTTTGGGACACTGGTTTGGCCATATTTGGCCCTTTGCTGATATCTGTGTGCTC 1631  
Qy 1253 TAAATCTTGTGTGTACAAATCTTGGCCGAATCTGTGAGTTCAGCCCAACTTCTCTTGTCT 1312  
Db 1632 TGAGTTTATTTGTGATCTTGTGTTTAAAGAAATGCCATTTGAACAC---CCAGTTC 1688  
Qy 1313 GTGTCAATGCTGTGCTGCTGCTTATACCGGAGAAAAAATGGTTTCATGAGCCGTCGGTTA 1372  
Db 1689 GAACCAATCAGATTCACAGTTCAGATTCCTGAACAGTCTGTACACGAGCCCTTGGCTG 1748  
Qy 1373 TTGTTTGGCTGGTGGAAATTTTACCTTTTGGTTTCAATCTTATTTGAATGATTTTCACTCT 1432  
Db 1749 GTATTATCATGGGAGGATTTTGGCCCTTTGGCTGCTGATCTTTTATACAACTTTTCTTCA 1808  
Qy 1433 TCAGCTCTTTCTGGGCATATAAGATCTATTATCTCTATGCTTTCATGATGCTGTGCTGG 1492  
Db 1809 TGAATAGTATTTGCTCACACAGATTTATTAATGATTTTGGCTTCTTATTTCTGGTGTTA 1868  
Qy 1493 TTATCTGTGATGTTGATCTGCTGTGTGATATTTGTGCAATATTTTCTACTAAATG 1552  
Db 1869 TCATTTTGGTTATTAATCTGTTCTGAAGCACTATATCTTCTGTCTATTTCCACCTATGTG 1928  
Qy 1553 CAGAAGATACCGTGGCATGCAAGTTTCTCTCTGCTGCTGATCACTCACTGCTATG 1612  
Db 1929 CAGAGGATTTATCATTTGGCAATGGCGTTCACTTCTTACGAGTGGCTTCTTCTTCTTAT 1988  
Qy 1613 TTTACATGATTTCTTTTACTACTATTTTTCATTTTCAAAACAAAGATGATGGCTTATTTCAA 1672  
Db 1989 TCTTAATCATGAGTACACTACTTCTTTTCAAACTGCAATGCACTGCAAGCA 2048  
Qy 1673 CATCATTTTACTTTGATATATGCGGTATTTAGCACAGCCTTTGGGGATTAATGTGTGAG 1732  
Db 2049 CAATTCGTACTTTGGTTATACCATGATGTTTGTGATCTTCTTTTCTTTTACAGGAA 2108  
Qy 1733 CGATTGTTACATGGGAACAGTGGCTTTGTCGGAATAATCTATCTAATGTGAAATTTG 1792  
Db 2109 CAATGGCTTCTTTGCTGATGCTTTTGGTTTGTACCAAAATATACAGTGTGGAAGTTG 2168  
Qy 1793 ACTAGAGA 1800  
Db 2169 ACTGAGA 2176

RESULT 12  
US-10-104-047-1699  
; Sequence 1699, Application US/10104047  
; Patent No. 6943241  
; GENERAL INFORMATION:  
; APPLICANT: HELIX RESEARCH INSTITUTE  
; TITLE OF INVENTION: No. 6943241el full length cDNA  
; FILE REFERENCE: H1-A0105  
; CURRENT APPLICATION NUMBER: US/10/104,047  
; CURRENT FILING DATE: 2002-03-25  
; PRIOR APPLICATION NUMBER:  
; PRIOR FILING DATE:  
; NUMBER OF SEQ ID NOS: 4096  
; SOFTWARE: PatentIn Ver. 2.1  
; SEQ ID NO 1699  
; LENGTH: 1878  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
US-10-104-047-1699

Query Match 10.9%; Score 226; DB 3; Length 1878;

[illegible]

Db	1698	CAATTCGTGACTTTGGTTATACCATGATGAATGGTTTGTGATCTCTCTTTTACAGGAA	1755
Qy	1733	CGATTGGTTACATGGGAAACAAGTCCTTTGTGCGAAAAATCTATATAAATGTGAAAAATTG	1792
Db	1758	CAATTGGCTCTCTTTGTATGCTTTTGGTTGTACCAAAATATACAGTGTGGTGAAGTTG	1817
Qy	1793	ACTAGAGA	1800
Db	1818	ACTGAAGA	1825

RESULT 13  
 US-09-270-767-14715  
 ; Sequence 14715, Application US/09270767  
 ; Patent No. 6703491  
 ; GENERAL INFORMATION:  
 ; APPLICANT: Homburger et al.  
 ; TITLE OF INVENTION: Nucleic acids and proteins of *Drosophila melanogaster*  
 ; FILE REFERENCE: File Reference: 7326-094  
 ; CURRENT APPLICATION NUMBER: US/09/270,767  
 ; CURRENT FILING DATE: 1999-03-17  
 ; NUMBER OF SEQ ID NOS: 62517  
 ; SOFTWARE: PatentIn Ver. 2.0  
 ; SEQ ID NO 14715  
 ; LENGTH: 995  
 ; TYPE: DNA  
 ; ORGANISM: *Drosophila melanogaster*  
 US-09-270-767-14715

Qy	95	TGCCCCGGACCCGGGGGACGACGACGATATCAAGATTAAGAGGAGTGTCT	154
Db	532	TGTCACTCTCCAGGCAGATGAGCAATCACAAGTACAATGACCGGGAGGAGGTGTAC	591
Qy	155	TATGATGAATACTGTGGCCCTACCATAATCGTCAAGAAACATATAAGTACTTTTCAC	214
Db	592	TGTGATGAACACGGTGGGCCCGTACCACAACTCGGAGGAGACGTACCGGTACTTCTCTC	651
Qy	215	TTCCATTCTGTGTGGGTCAAAAAAAGTATCAGTCAATTACCATGAAACTCTGGGAGAAAG	274
Db	652	TCCCTTTTGAGTGGCCAAAGTCTCGATATCCACTACACGAGACGCTGAGCGAGG	711
Qy	275	CACTTCAAGGGGTGCAATTTGGAATTAGTGGTCTGGATATTAAATTTAAAGATGATGA	334
Db	712	CGCTGCAAGGAGTCGAGCTGGAGTTTCAGTGGCTACGAGATGAGTTCAAGAGCGACGCC	771
Qy	335	TGCCAGCCACTTACTGTGGAATTGATTTAGATAAAGAAAGAGAGATGCAATTTGTATATG	394
Db	772	CCAAATTCGGTCACTCTGCATGGTCCACTTGCAGGAGGAGCGCCCAAGGCATTCACCTATG	831
Qy	395	CCATAAAAAATCAATTACTGTGTACCAAGATGTACATAGATGATTTACCAATATGGGTTATG	454
Db	832	CCGTGAAGAACCGAGTACTGGTACCAAAATGTACATGATGGAATGCCCAATTTGGGGAAG	891
Qy	455	TTGGTGGAGCTCATGAAAATGCGAAGATTACTATCTTTTGGACCTATAAAAACTTGA	514
Db	892	TGGGTGAGCGACGACGCGGATGCGAGTACTATATCTTCAAGCAGAGAGTTCGACA	951
Qy	515	TAGGTTTAAATCGAATCGAATTTGTGATGTTAACTTA	556
Db	952	TGGGTACTAATGGCCAGCAAAATCGTGGATATCACCTGACCA	993

Query Match 7.8%; Score 161.2; DB 3; Length 995;  
 Best Local Similarity 59.3%; Pred. No. 1.1e-27;  
 Matches 274; Conservative 0; Mismatches 188; Indels 0; Gaps 0;

## RESULT 14

RECORD 14  
US-09-248-796A-6208  
; Sequence 6208, Application US/03248796A  
; Patent No. 6747137  
; GENERAL INFORMATION:  
; APPLICANT: Keith Weinstein et al  
; TITLE OF INVENTION: NUCLEIC ACID AND A

; TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS

; FILE REFERENCE: 107196.132

; CURRENT APPLICATION NUMBER: US/09/248,796A

; CURRENT FILING DATE: 1999-02-12

; PRIOR APPLICATION NUMBER: US 60/074,725

; PRIOR FILING DATE: 1998-02-13

; PRIOR APPLICATION NUMBER: US 60/096,409

; PRIOR FILING DATE: 1998-08-13

; NUMBER OF SEQ ID NOS: 28208

; SEQ ID NO 6208

; LENGTH: 726

; TYPE: DNA

; ORGANISM: Candida albicans

US-09-248-796A-6208

Query Match 6.4%; Score 132.8; DB 3; Length 726;

Best Local Similarity 51.0%; Pred. No. 3.8e-21;

Matches 367; Conservative 0; Mismatches 347; Indels 6; Gaps 2;

QY 1078 GGAGGAGGAGATGATGAAGAAGCAGATGTTTATTTGGGCAATCCTTATCCAGCTATGGTG 1137

Db 13 GGTGGTGACAAATGGAAATTCATATGTTTGTGACACAGTTTGTAGTACCAAGGATTTTG 72

QY 1138 TGTGGCACTGCTTCTTCATCAATTTTCATAGCCATTTATTACCATGCTTCAAGAGCCATT 1197

Db 73 TCTCTGGTTTCGTGTGTGAATTTCTTTTAATTCAGTACAACTCTTCTGGTGTATT 132

QY 1198 CTTTTCGGAACAATGGTGGCGGTTTGTGCATCTGTTTTTGTATCTTCCTCTCAAT 1257

Db 133 CATATGGGGAACAATGTTTGGCAATGTCTTAATTTGGTTCATTTATATGATTCATTAAAT 192

QY 1258 CTTGTTGCTACAACTACTTGGCGGAAATCTGTGAGTCAAGCCAACTTTCTTGTGCGTTC 1317

Db 193 GTTATTGATCAATTTTAGCTAGTAATAGACCATATTATTC---GGTACAGTGAGAACT 249

QY 1318 AATGCTGTGCTCTCTCTATACCGGAGAAATAATGGTTCATGAGAGCCTGCGGTTATTGTT 1377

Db 250 AATCAAAATCCAGACAAATTCCTACTCAACCATGGTATTTAAGTACTACTCCCGGTAATG 309

QY 1378 TGCCTGGGTGGAATTTTACCTTTTGGTTCATCTTTAATGAAATGTAATTCATCTTCACG 1437

Db 310 TTTATTTTCGGGAATTTTCCCAATTTGGATCAATTCCTGTGGAAATGTAATTTTATTATTC 369

QY 1438 TCTTTCTGGGCATATAGATCTATTATGCTATGCTTCAATGATGCTGGTGTGCTGTTATC 1497

Db 370 TCAATTTGGTTTAAAGATTTTATATGTTTGGATTTTATTTTCTGTTCATATTA 429

QY 1498 CTGTGCAATGTGACTGTCTGTGACTATTGTGTCACATATTTTCTACTAAATGCAAGAA 1557

Db 430 ATGATTTTAACACTAGTTTAAATTAATTTTAAATGATTTATTATCTTTTATGTTTCAGAA 489

QY 1558 GATTACCGGTGGCAATGGAACAGTTTCTCTGCTGATCACTGCAATCTATGTTTAC 1617

Db 490 AATTATAAATGGCAATGGAATCATTTATTTGTGGAGAGGTTGTGCAATTTATGTTATTT 549

QY 1618 ATGATTTCTTTTACTACTATTTTTCAAAACAAGATGATGCTTATTTCAACAATCA 1677

Db 550 ATTCATTCATTTTTTTTGAAGCTGGTGGTA---AAATTTGGTGAATTTAGTTTCAATGTT 606

QY 1678 TTTTACTTTTGGATATATGGCGGTATTTTAGCAGCCTTTGGGGATAATGTGTGGAGCGAAT 1737

Db 607 TTATACAGTGGTTATTCAGCTGTGATTTCAATTTATGTTTTCCTTTTGTGGATCAAT 666

QY 1738 GGTTCATGGGAACAGTGGCTTTGTCCGAAAATCTATCTATGTAATGAAAATGACTAG 1797

Db 667 GGATTTTATAGTATTTAATATTTTGTACAGATTAATTTTATGCTCAATTTAAATTTGATTG 726

RESULT 15

US-09-949-016-1721/c

; Sequence 1721, Application US/09949016

; Patent No. 6812339

; GENERAL INFORMATION:

; APPLICANT: VENTER, J. Craig et al.

; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED

; WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF

; FILE REFERENCE: CL001307

; CURRENT APPLICATION NUMBER: US/09/949,016

; CURRENT FILING DATE: 2000-04-14

; PRIOR APPLICATION NUMBER: 60/241,755

; PRIOR FILING DATE: 2000-10-20

; PRIOR APPLICATION NUMBER: 60/237,768

; PRIOR FILING DATE: 2000-10-03

; PRIOR APPLICATION NUMBER: 60/231,498

; PRIOR FILING DATE: 2000-09-08

; NUMBER OF SEQ ID NOS: 207012

; SOFTWARE: FastSeq for Windows Version 4.0

; SEQ ID NO 1721

; LENGTH: 499

; TYPE: DNA

; ORGANISM: Human

US-09-949-016-1721

Query Match 5.9%; Score 122; DB 3; Length 499;

Best Local Similarity 100.0%; Pred. No. 1.1e-18;

Matches 122; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1934 TTACGAATGAGGCAATTTATTAAACCCAGCATGGTTTCTTAATGCTTCTTGTGGCAGC 1993

Db 499 TTACGAATGAGGCAATTTATTAAACCCAGCATGGTTTCTTAATGCTTCTTGTGGCAGC 440

QY 1994 TGCCACCTGTCCGGCGAATCTGTCCAGATCTTTTGTCCCTGAGGTGTCAGTTTGGGCC 2053

Db 439 TGCCACCTGTCCGGCGAATCTGTCCAGATCTCTTTGTCCCTGAGGTGTCAGTTTGGGCC 380

QY 2054 GC 2055

Db 379 GC 378

Search completed: March 10, 2006, 21:55:57

Job time : 360.113 secs



Result No.	Query			Description	
	Score	Match	Length	ID	
1	2072	100.0	2072	8	US-10-755-466-1
2	1840.4	88.8	3508	3	US-09-814-353-11837
3	1840.4	88.8	4024	5	US-10-198-846-10005
4	1830.8	88.4	3370	3	US-09-374-046A-25
5	1830.8	88.4	3370	7	US-10-616-263-25
6	1814	87.5	1827	8	US-10-755-466-3
7	1789.4	86.4	3076	3	US-09-915-582-29
8	1789.4	86.4	3076	6	US-10-277-802-29
9	1789.4	86.4	3076	6	US-10-896-972-29
10	1749.4	84.4	3389	6	US-10-205-219-122
11	1749.4	84.4	3389	9	US-10-986-157-2297
12	1749.4	84.4	3389	9	US-10-287-436A-335
13	1051	50.7	6197	6	US-10-062-674-1697
14	749.6	36.2	1070	6	US-10-264-237-1414
15	656	31.7	1867	3	US-09-915-582-13
16	656	31.7	1867	6	US-10-277-802-13
17	656	31.7	1867	6	US-10-896-972-13
18	646	31.2	1863	10	US-11-097-143-22277
19	583.6	28.2	2461	8	US-10-435-115-140808
20	581	28.0	2355	8	US-10-719-930-4365
21	570.2	27.5	1899	7	US-10-437-963-39405
22	570	27.5	2406	7	US-10-437-963-14430
23	567.8	27.4	2698	8	US-10-435-115-140919



Db	830	TATGGATGGAAACAGGTGCGATGGAGATGTATTTAGACCATCAAGTCAACCCACTGATATTT	888
Qy	901	TCCTCTCTGATGGTCTCGGATCTCAGATTAATTTGCTGTGCTCTCATCGTTATTAATGTT	960
Db	890	TCCTCTCTGATGGTCTCGGATCTCAGATTAATTTGCTGTGCTCTCATCGTTATTAATGTT	949
Qy	961	GCAATGATGAAGATTAATATACGTAGAGGGGATCAATGCTCAGTACAGCCATATTGTC	1020
Db	950	GCAATGATGAAGATTAATATACGTAGAGGGGATCAATGCTCAGTACAGCCATATTGTC	1009
Qy	1021	TATGCTGCTACGTCTCCAGTGAATGGTATTTTGGAGGAAGTCTGTATGCTAGACAAGGA	1080
Db	1010	TATGCTGCTACGTCTCCAGTGAATGGTATTTTGGAGGAAGTCTGTATGCTAGACAAGGA	1069
Qy	1081	GGAAGGAGATGGATAAAGCAGATGTTTATTTGGGGCAATCTTATCCAGCTATGGTGTGT	1140
Db	1070	GGAAGGAGATGGATAAAGCAGATGTTTATTTGGGGCAATCTTATCCAGCTATGGTGTGT	1129
Qy	1141	GGCACTGGCTTCTTTCATCAATTTTCATAGCCATTTATACCATGCTTCAAGAGCCATTCCT	1200
Db	1130	GGCACTGGCTTCTTTCATCAATTTTCATAGCCATTTATACCATGCTTCAAGAGCCATTCCT	1189
Qy	1201	TTTGGAAACAATGCTGGCGGTTGCTGTCATCTGTTTTTTTGGTATTTCTTCTCTAAATCTTT	1260
Db	1190	TTTGGAAACAATGCTGGCGGTTGCTGTCATCTGTTTTTTTGGTATTTCTTCTCTAAATCTTT	1249
Qy	1261	GTTGGTCAATACTTGGCCGAAATCTGTCAAGTTCAGGTCAGCCAACTTTCTTGTCTGTCAAT	1320
Db	1250	GTTGGTCAATACTTGGCCGAAATCTGTCAAGTTCAGGTCAGCCAACTTTCTTGTCTGTCAAT	1309
Qy	1321	GCTGTGCTGCTCTATACCGGAGAAAAATGTTTCATGAGGCTCGGGTTATTTGTTGC	1380
Db	1310	GCTGTGCTGCTCTATACCGGAGAAAAATGTTTCATGAGGCTCGGGTTATTTGTTGC	1369
Qy	1381	CTGGGTGGAAATTTTACCTTTTGGTTCAACTCTTATTTGAAATGTATTTCACTTCACTCT	1440
Db	1370	CTGGGTGGAAATTTTACCTTTTGGTTCAACTCTTATTTGAAATGTATTTCACTTCACTCT	1429
Qy	1441	TTCTGGGCATATAAGATCTATTATGTCTATGGCTTCATGATGCTGTGTGTTATCTCTG	1500
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Qy	1501	TGCATTTGCACTGCTGTGTGACTATTTGTGTGCACATATTTTCTCTATTAATGACAGAGAT	1560
Db	1490	TGCATTTGCACTGCTGTGTGACTATTTGTGTGCACATATTTTCTCTATTAATGACAGAGAT	1549
Qy	1561	TACCGTGGCAATGGACAAGTTTCTCTCTGCTGCATCAACTGCAATCTATGTTTACATG	1620
Db	1550	TACCGTGGCAATGGACAAGTTTCTCTCTGCTGCATCAACTGCAATCTATGTTTACATG	1609
Qy	1621	TATTTCTTTTACTACTATTTTTTCAAAACAAGATGTATGGCTTATTTCAAAACATCATTT	1680
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Qy	1681	TACTTTGGATATATANGCGGTATTTAGCACAGCTTGGGGATTAATGTGTGGAGCGATTGGT	1740
Db	1670	TACTTTGGATATATANGCGGTATTTAGCACAGCTTGGGGATTAATGTGTGGAGCGATTGGT	1729
Qy	1741	TACATGGGACAAGTGCCTTTGTCGAAAAATCTATCTAAATCTGAAAAATTGACTAGAGA	1800
Db	1730	TACATGGGACAAGTGCCTTTGTCGAAAAATCTATCTAAATCTGAAAAATTGACTAGAGA	1789
Qy	1801	CCCAAGAAAACCTGGAACTTTTGGATCAAATTTCTTTTTCATAGGGGTGGAACCTTGCACAGC	1860
Db	1790	CCCAAGAAAACCTGGAACTTTTGGATCAAATTTCTTTTTCATAGGGGTGGAACCTTGCACAGC	1849
Qy	1861	AAAAACAACAAACGACAAGAGATTGGGCTTTAACTTTTTTTTTTTTTTTTTTTTTTTTTT	1920
Db	1850	AAAAACAACAAACGACAAGAGATTGGGCTTTAACTTTTTTTTTTTTTTTTTTTTTTTTTT	1909
Qy	1921	TTTTTTT 1926	
Db	1910	CTCTTT 1915	













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QY 982 ACTGAGAGGGGATCAATGCTCAGTACAGCCATATTTGCTATGCTGCTAGCTCCAGTG 1041
DB 913 ACTGAGAGGGGATCAATGCTCAGTACAGCCATATTTGCTATGCTGCTAGCTCCAGTG 972
QY 1042 AATGGTTATTTTGAGGAAGTCTGTATGCTAGACAGGAGGAGAGATGATTAAGCAG 1101
DB 973 AATGGTTATTTTGAGGAAGTCTGTATGCTAGACAGGAGGAGAGATGATTAAGCAG 1032
QY 1102 ATGTTTATTTGGGGCATTCCTTATCCAGCTATGCTGTGCTGCTGCTTCTTCTCATCAAT 1161
DB 1033 ATGTTTATTTGGGGCATTCCTTATCCAGCTATGCTGTGCTGCTGCTTCTTCTCATCAAT 1092
QY 1162 TTCAATGCCATTTATACCATGCTTCAAGAGCCATTCCTTTTGGAACAATGGTGGCGGTT 1221
DB 1093 TTCAATGCCATTTATACCATGCTTCAAGAGCCATTCCTTTTGGAACAATGGTGGCGGTT 1152
QY 1222 TGTGTCATCTGTTTTTTGTTATTTCTTCTCTCAATCTGTTGTCATCAATCTTGGCCGA 1281
DB 1153 TGTGTCATCTGTTTTTTGTTATTTCTTCTCTCAATCTGTTGTCATCAATCTTGGCCGA 1212
QY 1282 AATCTGTGAGTCCAGCCCACTTCTTGTGCTGCTCAATGCTGCTGCTGCTTCTTCTATACCG 1341
DB 1213 AATCTGTGAGTCCAGCCCACTTCTTGTGCTGCTCAATGCTGCTGCTGCTTCTTCTATACCG 1272
QY 1342 GAGAAAAATGGTTCATGAGGAGCCGCGGTTATTTGTTGCTGCTGCTGCTGCTGCTGCTGCT 1401
DB 1273 GAGAAAAATGGTTCATGAGGAGCCGCGGTTATTTGTTGCTGCTGCTGCTGCTGCTGCTGCT 1332
QY 1402 GGTTCATCTTTATGAAATGATTTCTCATCTTCTCAAGTCTTCTGCTGCTGCTGCTGCTGCT 1461
DB 1333 GGTTCATCTTTATGAAATGATTTCTCATCTTCTCAAGTCTTCTGCTGCTGCTGCTGCTGCT 1392
QY 1462 TATGCTATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1521
DB 1393 TATGCTATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1452
QY 1522 ACTATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1581
DB 1453 ACTATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1512
QY 1582 TTTCTCTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1641
DB 1513 TTTCTCTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1572
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DB 1573 TTCAAAAAAGAGATGATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1632
QY 1702 TTTAGCACACCTTTGGGGAATATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1761
DB 1633 TTTAGCACACCTTTGGGGAATATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1692
QY 1762 GTCCGAAAAATCTATCTATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1821
DB 1693 GTCCGAAAAATCTATCTATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1752
QY 1822 GGATCAATTTCTTTTTCATAGGGGTGGAATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1881
DB 1753 GGATCAATTTCTTTTTCATAGGGGTGGAATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1812
QY 1882 GAGATTGGGCTTTAACTTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTT 1926
DB 1813 GAGATTGGGCTTTAACTTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTT 1857

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RESULT 8

US-10-277-802-29  
; Sequence 29, Application US/10277802  
; Publication No. US2003019070A1  
; GENERAL INFORMATION:  
; APPLICANT: Rosen et al.  
; TITLE OF INVENTION: 17 Human Secreted Proteins

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; FILE REFERENCE: PS723P1
; CURRENT APPLICATION NUMBER: US/10/277,802
; CURRENT FILING DATE: 2002-10-23
; PRIOR APPLICATION NUMBER: 09/915,582
; PRIOR FILING DATE: 2001-07-27
; PRIOR APPLICATION NUMBER: PCT/US01/01431
; PRIOR FILING DATE: 2001-01-17
; PRIOR APPLICATION NUMBER: 60/179,065
; PRIOR FILING DATE: 2000-01-31
; PRIOR APPLICATION NUMBER: 60/180,628
; PRIOR FILING DATE: 2000-02-04
; PRIOR APPLICATION NUMBER: 60/231,968
; PRIOR FILING DATE: 2000-09-12
; NUMBER OF SEQ ID NOS: 97
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 29
; LENGTH: 3076
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (3064)
; OTHER INFORMATION: n equals a,t,g, or c
US-10-277-802-29

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Query Match 86.4%; Score 1789.4; DB 6; Length 3076;  
Best Local Similarity 98.0%; Pred. No. 0;  
Matches 1808; Conservative 3; Mismatches 34; Indels 0; Gaps 0;

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QY 82 CTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 141
DB 13 CTGAGGTTACCGTCCGGAATTCCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 72
QY 142 GAGGAAGTTGTTCTTATGATGAATCTGTTGGGCCCTACCAATAATCGTCAAGAAACATAT 201
DB 73 GAGGAAGTTGTTCTTATGATGAATCTGTTGGGCCCTACCAATAATCGTCAAGAAACATAT 132
QY 202 AAGTACTTTTCACTTCCATCTGTTGGGTCCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 261
DB 133 AAGTACTTTTCACTTCCATCTGTTGGGTCCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 192
QY 262 ACTCTGGGAGAGACACTTCAAGGGTTGAAATTTAGTGGTCTGCTGCTGCTGCTGCTGCTGCT 321
DB 193 ACTCTGGGAGAGACACTTCAAGGGTTGAAATTTAGTGGTCTGCTGCTGCTGCTGCTGCTGCT 252
QY 322 AAGATGATGCTGATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 381
DB 253 AAGATGATGCTGATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 312
QY 382 GAATTTGATATGCCATAAAAAATCACTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 441
DB 313 GCATTTGTATATGCCATAAAAAATCACTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 372
QY 442 ATATGGGGTATTTGTTGGTGGCTGATGAAATGGAAGAGATTTACTATCTTTGGACCTAT 501
DB 373 ATATGGGGTATTTGTTGGTGGCTGATGAAATGGAAGAGATTTACTATCTTTGGACCTAT 432
QY 502 AAAAAAATTGAAATAGTTTTTAATGGAATTCGAAATTTGTTGATGTTAATCTAACTAGTGA 561
DB 433 AAAAAAATTGAAATAGTTTTTAATGGAATTCGAAATTTGTTGATGTTAATCTAACTAGTGA 492
QY 562 GGAAGAGTGAATCTGTTGCCAATAATCACTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 621
DB 493 GGAAGAGTGAATCTGTTGCCAATAATCACTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 552
QY 622 AAGTCAGATGCTGAAATTTGGAAGATCGATTTGCAAAATATCTTGATCCGCTCTTTTTCCT 681
DB 553 AAGTCAGATGCTGAAATTTGGAAGATCGATTTGCAAAATATCTTGATCCGCTCTTTTTCCT 612
QY 682 CATCGGATTCATTTGGTTTTTCAATTTTCAATCTCTTCATGATGGTGTATCTTTTGGGGC 741
DB 613 CATCGGATTCATTTGGTTTTTCAATTTTCAATCTCTTCATGATGGTGTATCTTTTGGGGC 672

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110 GAAACACAGTATCAGATTAAGAGGAGTTGCTTAAGATGAATCTGTTGGCCCTAC 169  
181 CATATCGTCAAGAAACATATAAGTACTTTTCACTTCCATCTGTGTGGGGTCAAAAAA 240  
170 CATATCGTCAAGAAACATATAAGTACTTTTCACTTCCATCTGTGTGGGGTCAAAAAA 229  
241 AGTATCAGTCAATACCATCAATCTCGGAGAGCACTTCAAGGGGTGAATTCGAATTT 300  
230 AGTATCAGTCAATACCATCAATCTCGGAGAGCACTTCAAGGGGTGAATTCGAATTT 289  
301 AGTGTCTCGATATAAATTTAAAGATGATGTGATGCCATAAAAAATCATTTCTGGTACCG 360  
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361 TTAGATAAAGAAAGAGAGATGATTTGATATGCCATAAAAAATCATTTCTGGTACCG 420  
350 TTAGATAAAGAAAGAGAGATGATTTGATATGCCATAAAAAATCATTTCTGGTACCG 409  
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766 TTATGCTCGTACAGTAAAGAGAGAAATGATGATGATGATGATGATGATGATGATGATG 825  
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946 TGCATGATGATGATTTATATACGAGAGGATCAATGCTCAGTACAGCCATTTTGT 1005  
1020 CTATGCTGCTGCTCTCCAGTGAATGGTTTATTTGGAGGAGTCTGTATGCTAGACAAAG 1079  
1006 CTATGCTGCTGCTCTCCAGTGAATGGTTTATTTGGAGGAGTCTGTATGCTAGACAAAG 1065  
1080 AGGAAGGAGATGATGATGATGATTTATTTGGGAGATTCCTATCCAGCTATGCTGTG 1139  
1066 AGGAAGGAGATGATGATGATGATTTATTTGGGAGATTCCTATCCAGCTATG--GGT 1123  
1140 TGGCACTGCTCTCTCATCAATTTTCAAGCATTTTATACCATGCTTCAAGAGCCATTC 1199  
1124 GTGCACTGCTCTCTCATCAATTTTATAGCCATTTTATACCATGCTTCAAGAGCCATTC 1183  
1200 TTTTGGAAACATGTTGGGGCTTTGTTGATCTGTTTTTTTGTATTTCTCTCTAAATCT 1259

1184 TTTTGGAAACATGTTGGGGCTTTGTTGATCTGTTTTTTTGTATTTCTCTCTAAATCT 1243  
1260 TTTTGGTACAAATCTTGGCCGAAATCTGTGAGGTGAGCCCAACATTTCTTGTCTGTGCA 1319  
1244 TTTTGGTACAAATCTTGGCCGAAATCTGTGAGGTGAGCCCAACATTTCTTGTCTGTGCA 1303  
1320 TGTGTCCTGCTGCTATACCGGAGAAATGCTTCAATGAGCTCGGTTATTTGTTG 1379  
1304 TGTGTCCTGCTGCTATACCGGAGAAATGCTTCAATGAG--CTGGGTTATTTGTTG 1362  
1380 CTTGGTGGAAATTTTACCTTTTGGTTCAATCTTTTATGAAATGATTTTCACTTCAAGTC 1439  
1363 CTTGGTGGAAATTTTACCTTTTGGTTCAATCTTTTATGAAATGATTTTCACTTCAAGTC 1422  
1440 TTTTCTGGGCATATAAGATCTATTTATGCTATGCTTCAATGAGTGTGCTGCTGTTATCCT 1499  
1423 TTTTCTGGGCATATAAGATCTATTTATGCTATGCTTCAATGAGTGTGCTGCTGTTATCCT 1482  
1500 GTGCATTTGCTGCTGCTGCTGCTATTTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1559  
1483 GTGCATTTGCTGCTGCTGCTGCTATTTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1542  
1560 TTAACGGTGGCAATGGAAGTCTTCTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1619  
1543 TTAACGGTGGCAATGGAAGTCTTCTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1602  
1620 GTATTCCTTTTACTACTATTTTTCATGAAACAAAGATGATGCTTATTTTCAACATCATTT 1679  
1603 GTATTCCTTTTACTACTATTTTTCATGAAACAAAGATGATGCTTATTTTCAACATCATTT 1662  
1680 TTAACGGTGGCAATGGAAGTCTTCTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1739  
1663 TTAACGGTGGCAATGGAAGTCTTCTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1722  
1740 TTAACGGTGGCAATGGAAGTCTTCTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1799  
1723 TTAACGGTGGCAATGGAAGTCTTCTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1782  
1800 ACCCAAGAAACCTGGAACCTTGGATCAATTTCTTTTCTAGGGTGGAGCTTGCACAG 1859  
1783 ACCCAAGAAACCTGGAACCTTGGATCAATTTCTTTTCTAGGGTGGAGCTTGCACAG 1842  
1860 CAAAAACAAACAAACGCAAGAGATTTGGGCTTTAACTTTTTTTTTTTTTTTTTTTTTT 1919  
1843 CAAAAACAAACAAACGCAAGAGATTTGGGCTTTAAACACATGSGTACTTTTGGGTC 1902  
1920 TTTTCTTT 1926  
1903 TCTCTTT 1909

## RESULT 13

US-10-062-674-1697  
; Sequence 1697, Application US/10062674  
; Publication No. US20040005559A1  
; GENERAL INFORMATION:  
; APPLICANT: Loring, Jeanne P.; Kaser, Matthew R.  
; TITLE OF INVENTION: MARKERS OF NEURONAL DIFFERENTIATION AND MORPHOGENESIS  
; FILE REFERENCE: PA-0026-1 CIP  
; CURRENT APPLICATION NUMBER: US/10/062,674  
; CURRENT FILING DATE: 2002-01-30  
; PRIOR APPLICATION NUMBER: US 09/625,102  
; PRIOR FILING DATE: 2000-07-24  
; NUMBER OF SEQ ID NOS: 2217  
; SOFTWARE: PERL Program  
; SEQ ID NO 1697  
; LENGTH: 6197  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; FEATURE:  
; NAME/KEY: misc feature  
; OTHER INFORMATION: Incyte ID No. US20040005559A1 233927.4  
; FEATURE:





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; PRIOR FILING DATE: 2000-02-04
; PRIOR APPLICATION NUMBER: 60/231,968
; PRIOR FILING DATE: 2000-09-12
; NUMBER OF SEQ ID NOS: 97
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 13
; LENGTH: 1867
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-915-582-13

Query Match      31.7%; Score 656; DB 3; Length 1867;
Best Local Similarity 97.1%; Pred No. 5, 7e-140; Indels 0; Gaps 0;
Matches 668; Conservative 0; Mismatches 20;

Qy 1239 TGTATTCTTCCTCTAAATCTTGTTGGTACAAATCTTGGCCGAAATCTGTCAAGTCAAGCC 1298
Db      |||
19 TGTGCTCTTCTCTAAATCTTGTTGGTACAAATCTTGGCCGAAATCTGTCAAGTCAAGCC 78

Qy 1299 CAACCTTTCCTTGTCTGTCAATGCTGTGCTCTCTATACCGGAGAAATGTTTCAT 1358
Db      |||
79 CAACCTTTCCTTGTCTGTCAATGCTGTGCTCTCTATACCGGAGAAATGTTTCAT 138

Qy 1359 GGAGCCTGCGGTATATGTTTCCCTGGGTGGAAATTTACCTTTTGGTCAATCTTTATTGA 1418
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139 GGAGCCTGCGGTATATGTTTCCCTGGGTGGAAATTTACCTTTTGGTCAATCTTTATTGA 198

Qy 1419 AATGTAATTTCAATCTTACGTCCTTTCTGGGCATATAAGATCTATTATGTCTATGGCTTCAT 1478
Db      |||
199 AATGTAATTTCAATCTTACGTCCTTTCTGGGCATATAAGATCTATTATGTCTATGGCTTCAT 258

Qy 1479 GATGCTGGTGTGCTTATCTCTGTCATGTCACGTCTGTGTGACTATTGTGTGCACATA 1538
Db      |||
259 GATGCTGGTGTGCTTATCTCTGTCATGTCACGTCTGTGTGACTATTGTGTGCACATA 318

Qy 1539 TTTTCTACTAAATCAGAGATTAACCGTGGCAATGGACAAAGTTTCTCTCTGCTGCATC 1598
Db      |||
319 TTTTCTACTAAATCAGAGATTAACCGTGGCAATGGACAAAGTTTCTCTCTGCTGCATC 378

Qy 1599 AACTGCAATCTATCTTTACATGTAATCTTTTACTACTATTTTTCAAAAAAGATGTA 1658
Db      |||
379 AACTGCAATCTATCTTTACATGTAATCTTTTACTACTATTTTTCAAAAAAGATGTA 438

Qy 1659 TGGCTTATTTCAACATCATTTTACTTTGGATATATGCGGTATTTAGCACAGCCTTGGG 1718
Db      |||
439 TGGCTTATTTCAACATCATTTTACTTTGGATATATGCGGTATTTAGCACAGCCTTGGG 498

Qy 1719 GATAATGTGTGAGCGATTGGTTACATGGGAACAAGTGCCCTTTGTCGAAAAATCTATAC 1778
Db      |||
499 GATAATGTGTGAGCGATTGGTTACATGGGAACAAGTGCCCTTTGTCGAAAAATCTATAC 558

Qy 1779 TAATGTGAAAAATTCAGTAGAGACCCAAAGAAACCTGGAACCTTTCGATCAATTTCTTTTC 1838
Db      |||
559 TAATGTGAAAAATTCAGTAGAGACCCAAAGAAACCTGGAACCTTTCGATCAATTTCTTTTC 618

Qy 1839 ATAGGGGTGGNACTTGCACAGCAAAACAAACAAACAGAGAGATTTGGGCTTTAAC 1898
Db      |||
619 ATAGGGGTGGNACTTGCACAGCAAAACAAACAAACAGAGAGATTTGGGCTTTAAC 678

Qy 1899 TTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTTTCTTTT 1926
Db      |||
679 AACTGGGTACTTTTGGGTCTCTCTTT 706
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Job time : 1704.01 secs

GenCore version 5.1.7  
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OM nucleic - nucleic search, using sw model

Run on: March 10, 2006, 22:05:49 ; Search time 664.273 Seconds  
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Perfect score: 2072

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Searched: 7673375 seqs, 1153648444 residues

Total number of hits satisfying chosen parameters: 15346750

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : Published Applications NA New:\*

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13: /cgn2\_6/ptodata/2/pubna/US60\_NEW\_PUB.seq.\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	482.8	23.3	570	12	US-11-128-061-1325 Sequence 1325, Ap
2	482.8	23.3	570	12	US-11-128-061-1325 Sequence 4967, Ap
3	482.8	23.3	570	12	US-11-128-049-1325 Sequence 1325, Ap
4	482.8	23.3	570	12	US-11-128-049-1325 Sequence 4967, Ap
5	227.6	11.0	2391	12	US-11-000-688-1309 Sequence 1309, Ap
6	226	10.9	1878	9	US-11-072-512-1699 Sequence 1699, Ap
7	190.4	9.2	1094	9	US-11-240-769-50 Sequence 50, Appl
8	172	8.3	1821	9	US-11-240-769-49 Sequence 49, Appl
9	169.4	8.2	1816	9	US-11-240-769-21 Sequence 21, Appl
10	145	7.0	569	12	US-11-000-688-313 Sequence 313, Ap
11	138	6.7	1251	7	US-09-925-065A-724812 Sequence 724812, Ap
12	124.4	6.0	2019	7	US-10-932-182A-478 Sequence 478, Ap
13	124.4	6.0	2019	7	US-10-932-182A-478 Sequence 478, Ap
14	120	5.8	2025	7	US-10-932-182A-4690 Sequence 4690, Ap
15	120	5.8	2025	7	US-10-932-182A-4690 Sequence 4690, Ap
16	112.2	5.4	495	12	US-11-128-061-345 Sequence 345, Ap
17	112.2	5.4	495	12	US-11-128-061-3987 Sequence 3987, Ap
18	112.2	5.4	495	12	US-11-128-049-345 Sequence 345, Ap
19	112.2	5.4	495	12	US-11-128-049-3987 Sequence 3987, Ap
20	107.8	5.2	476	12	US-11-000-688-312 Sequence 312, Ap

21	101.6	4.9	424	12	US-11-000-688-1308 Sequence 1308, Ap
22	98.8	4.8	636	7	US-10-932-182A-173799 Sequence 173799, Ap
23	98.8	4.8	636	7	US-10-932-182A-173799 Sequence 173799, Ap
24	96	4.6	623	6	US-09-925-065A-799365 Sequence 799365, Ap
25	69.8	3.4	465	7	US-10-932-182A-173463 Sequence 173463, Ap
26	69.8	3.4	465	7	US-10-932-182A-173463 Sequence 173463, Ap
27	60.4	2.9	1083	7	US-10-932-182A-173986 Sequence 173986, Ap
28	60.4	2.9	1083	7	US-10-932-182A-173986 Sequence 173986, Ap
29	59.6	2.9	427	8	US-10-821-234-288 Sequence 288, App
30	53.8	2.6	1488	9	US-11-096-568A-25839 Sequence 25839, A
31	53.2	2.6	1478	8	US-10-909-125-1744 Sequence 1744, Ap
32	52.6	2.5	384	7	US-10-932-182A-81876 Sequence 81876, A
33	52.6	2.5	384	7	US-10-932-182A-81876 Sequence 81876, A
34	52.2	2.5	516	6	US-09-925-065A-480255 Sequence 480255, A
35	52	2.5	687	8	US-10-986-501-107 Sequence 107, App
36	50.6	2.4	486	6	US-09-925-065A-134718 Sequence 134718, Ap
37	50	2.4	704	6	US-09-925-065A-924411 Sequence 924411, Ap
38	49.8	2.4	106	8	US-10-310-914A-3758 Sequence 3758, Ap
39	49.8	2.4	505	6	US-09-925-065A-736759 Sequence 736759, Ap
40	49.8	2.4	505	6	US-09-925-065A-736760 Sequence 736760, Ap
41	49.8	2.4	627	6	US-09-925-065A-488892 Sequence 488892, Ap
42	49.8	2.4	182190	12	US-11-121-086-102 Sequence 102, App
43	49.6	2.4	534	6	US-09-925-065A-474669 Sequence 474669, Ap
44	49	2.4	486	6	US-09-925-065A-134717 Sequence 134717, Ap
45	49	2.4	486	6	US-09-925-065A-134719 Sequence 134719, Ap

#### ALIGNMENTS

##### RESULT 1

US-11-128-061-1325  
Sequence 1325, Application US/11128061  
Publication No. US20060003958A1  
GENERAL INFORMATION:  
APPLICANT: Melville, Mark W.  
APPLICANT: Charlebois, Timothy S.  
APPLICANT: Mounts, William M.  
APPLICANT: Hann, Louane E.  
APPLICANT: Sinacore, Martin S.  
APPLICANT: Leonard, Mark W.  
APPLICANT: Brown, Eugene L.  
APPLICANT: Miller, Christopher P.  
TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES RELATED TO OLIGONUCLEOTIDE ARRAYS  
TITLE OF INVENTION: TO MONITOR GENE EXPRESSION  
FILE REFERENCE: 01997.027701  
CURRENT APPLICATION NUMBER: US/11/128,061  
CURRENT FILING DATE: 2005-05-11  
PRIOR APPLICATION NUMBER: US 60/570,425  
PRIOR FILING DATE: 2004-05-11  
NUMBER OF SEQ ID NOS: 7285  
SOFTWARE: Patent in version 3.3  
SEQ ID NO 1325  
LENGTH: 570  
TYPE: DNA  
ORGANISM: Cricetus griseus

Query Match 23.3%; Score 482.8; DB 12; Length 570;

Best Local Similarity 91.6%; Pred. No. 5.8e-64;

Matches 522; Conservative 0; Mismatches 47; Indels 1; Gaps 1;

Qy 910 ATTGGTTCTGGAGTGCAGATATTGCTGCTCTCTCATCGTTATTATTGTCATGATATA 969  
Db 1 ATTGGTTCTGGAGTGCAGATATTGCTGCTCTCTCATCGTTATTATTGTCATGATATA 60

Qy 970 GAGATTATATATCTCAGAGGGGATCAATGCTCAGTACAGCCATATTGTCATGCTGCT 1029  
Db 61 GAGATTATATATACAGAGGGGATCAATGCTCAGTACAGCCATATTGTCATGCTGCT 120

Qy 1030 ACCTCTCCAGTGAATGGTTATTATTTGGAGGAAGTCTGTATGCTAGACAAGGAGGAGA 1089  
Db 121 ACATCTCCAGTGAATGGTTATTATTTGGAGGAAGTCTGTATGCTAGACAAGGAGGAGA 180



QY 1090 TGGATAAAGCAGATGTTTATTTGGGGCATTCCTTATCCAGCTATAGTGTGTGGCACTGCC 1149  
DB 181 TGGATAAAGCAGATGTTTATTTGGGGCATTCCTTATCCAGCTATAGTGTGTGGCACTGCC 240  
QY 1150 TTCTTCATCAATTTTCATAGCCATTTATACCATGCTTCAAGAGCCATTCCTTTTGGAA 1209  
DB 241 TTCTTCATCAATTTATAGCCATTTATATCATGCTCTAGAGCCATTCCTTTTGGAA 300  
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QY 1270 ATACTTGGCGGAATCTGTAGCTAGCCCAATCTTCTGTGCTGAATGCTGTGCT 1329  
DB 361 ATACTTGGCGGAATCTGTAGCTAGCCCAATCTTCTGTGCTGAATGCTGTGCT 420  
QY 1330 CGTCTATACCGAGAAAAATGGTTCATGGAGCCTGCGGTTATTTGCTTGGCTGGA 1389  
DB 421 GGTCTATCCAGACAAATATGGTTATGGATCCCTGCAATTTATCGCTGCTAGCA 480  
QY 1390 ATTTTACCTTTTGGTTCAATCTTTATTTGAATGATTTTCACTTCACTGCTTTCTGGGA 1449  
DB 481 ATTTTACCAATTTGGCTCCATCTTCAATGAAATGACTTTCATGTAACATCTTTCTGGGA 540  
QY 1450 TATAAGATCTA-TTATGCTATGGCTTCAT 1478  
DB 541 TACAAGACCCACTTATGCTATGGCTTTAT 570

## RESULT 2

US-11-128-061-4967  
; Sequence 4967, Application US/11128061  
; Publication No. US2006003958A1

## GENERAL INFORMATION:

; APPLICANT: Melville, Mark W.  
; APPLICANT: Charlebois, Timothy S.  
; APPLICANT: Mounts, William M.  
; APPLICANT: Hann, Louane E.  
; APPLICANT: Sinacore, Martin S.  
; APPLICANT: Leonard, Mark W.  
; APPLICANT: Brown, Eugene L.  
; APPLICANT: Miller, Christopher P.  
; TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES RELATED TO OLIGONUCLEOTIDE ARRAYS  
; FILE REFERENCE: 01997.027701  
; CURRENT APPLICATION NUMBER: US/11/128,061  
; PRIOR FILING DATE: 2005-05-11  
; PRIOR APPLICATION NUMBER: US 60/570,425  
; NUMBER OF SEQ ID NOS: 7285  
; SOFTWARE: Patent in version 3.3  
; SEQ ID NO 4967  
; LENGTH: 570  
; TYPE: DNA  
; ORGANISM: Cricetulus griseus

US-11-128-061-4967

Query Match 23.3%; Score 482.8; DB 12; Length 570;  
Best Local Similarity 91.6%; Pred. No. 5.8e-64;  
Matches 522; Conservative 0; Mismatches 47; Indels 1; Gaps 1;

QY 910 ATTGGTTCGGATGTCAGATATTTGCTGTGCTCTCATGTTATTTATTTGCAATGATA 969  
DB 1 ATTGGTTCGGATGTCAGATATTTGCTGTGCTCTCATGTTATTTATTTGCAATGATA 60  
QY 970 GAAGATTTATATCTAGAGGGGATCAATGCTCAGTACAGCCATATTTCTATGCTGCT 1029  
DB 61 GAGGATTTATATACAGAGAGGGGATCAATGCTCAGTACAGCCATATTTCTATGCTGCT 120  
QY 1030 ACCTCTCCAGTGAATGTTATTTTGGAGGAAGTCTGTATGCTAGACAGAGGAGGAGA 1089  
DB 121 ACATCTCCAGTGAATGTTATTTTGGAGGAAGTCTGTATGCTAGACAGAGGAGGAGA 180

QY 1090 TGGATAAAGCAGATGTTTATTTGGGGCATTCCTTATCCAGCTATAGTGTGTGGCACTGCC 1149  
DB 181 TGGATAAAGCAGATGTTTATTTGGGGCATTCCTTATCCAGCTATAGTGTGTGGCACTGCC 240  
QY 1150 TTCTTCATCAATTTTCATAGCCATTTATACCATGCTTCAAGAGCCATTCCTTTTGGAA 1209  
DB 241 TTCTTCATCAATTTATAGCCATTTATATCATGCTCTAGAGCCATTCCTTTTGGAA 300  
QY 1210 ATGGTGGCGGTTTGTGTCATCTGTTTTTTTGTATTTCTTAAATCTTGTGTGTA 1269  
DB 301 ATGGTGGCGGTTTGTGTCATCTGTTTTTTTGTATTTCTTAAATCTTGTGTGTA 360  
QY 1270 ATACTTGGCGGAATCTGTAGCTAGCCCAATCTTCTGTGCTGAATGCTGTGCT 1329  
DB 361 ATACTTGGCGGAATCTGTAGCTAGCCCAATCTTCTGTGCTGAATGCTGTGCT 420  
QY 1330 CGTCTATACCGAGAAAAATGGTTCATGGAGCCTGCGGTTATTTGCTTGGCTGGA 1389  
DB 421 GGTCTATCCAGACAAATATGGTTATGGATCCCTGCAATTTATCGCTGCTAGCA 480  
QY 1390 ATTTTACCTTTTGGTTCAATCTTTATTTGAATGATTTTCACTTCACTGCTTTCTGGGA 1449  
DB 481 ATTTTACCAATTTGGCTCCATCTTCAATGAAATGACTTTCATGTAACATCTTTCTGGGA 540  
QY 1450 TATAAGATCTA-TTATGCTATGGCTTCAT 1478  
DB 541 TACAAGACCCACTTATGCTATGGCTTTAT 570

## RESULT 3

US-11-128-049-1325  
; Sequence 1325, Application US/11128049  
; Publication No. US20060010513A1

## GENERAL INFORMATION:

; APPLICANT: Melville, Mark W.  
; APPLICANT: Charlebois, Timothy S.  
; APPLICANT: Mounts, William M.  
; APPLICANT: Hann, Louane E.  
; APPLICANT: Sinacore, Martin S.  
; APPLICANT: Leonard, Mark W.  
; APPLICANT: Brown, Eugene L.  
; APPLICANT: Miller, Christopher P.  
; TITLE OF INVENTION: OLIGONUCLEOTIDE ARRAYS TO MONITOR GENE EXPRESSION AND METHODS FOR  
; FILE REFERENCE: 01997.027700  
; CURRENT APPLICATION NUMBER: US/11/128,049  
; CURRENT FILING DATE: 2005-05-11  
; PRIOR APPLICATION NUMBER: US 60/570,425  
; PRIOR FILING DATE: 2004-05-11  
; NUMBER OF SEQ ID NOS: 7285  
; SOFTWARE: Patent in version 3.3  
; SEQ ID NO 1325  
; LENGTH: 570  
; TYPE: DNA  
; ORGANISM: Cricetulus griseus

US-11-128-049-1325

Query Match 23.3%; Score 482.8; DB 12; Length 570;  
Best Local Similarity 91.6%; Pred. No. 5.8e-64;  
Matches 522; Conservative 0; Mismatches 47; Indels 1; Gaps 1;

QY 910 ATTGGTTCGGATGTCAGATATTTGCTGTGCTCTCATGTTATTTATTTGCAATGATA 969  
DB 1 ATTGGTTCGGATGTCAGATATTTGCTGTGCTCTCATGTTATTTATTTGCAATGATA 60  
QY 970 GAAGATTTATATCTAGAGGGGATCAATGCTCAGTACAGCCATATTTCTATGCTGCT 1029  
DB 61 GAGGATTTATATACAGAGAGGGGATCAATGCTCAGTACAGCCATATTTCTATGCTGCT 120  
QY 1030 ACCTCTCCAGTGAATGTTATTTTGGAGGAAGTCTGTATGCTAGACAGAGGAGGAGA 1089  
DB 121 ACATCTCCAGTGAATGTTATTTTGGAGGAAGTCTGTATGCTAGACAGAGGAGGAGA 180



Db 1116 AAGATATTGCTAGATATATATCATGATGGACTTACGGAAGATGCCAG-----G 1163  
Qy 836 ATGAATATGGATGGAACACAGGTGCATGAGATGATATTTAGACCATCAAGTCAACCACTGA 895  
Db 1164 AAGAAATTTGGCTGGAACATTTGTTCTGATGATATTTCCGCTCCCAAGAAAGGATGC 1223  
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Db 1344 TGGTCTGCTGGGTGCTGCTGGGCACTCCCTGAGGCTATGTTGCTGCCAGATCTATAAGT 1403  
Qy 1073 GACAGGAGGAGGATGATAGGATAGGATGTTTATTTGGGCATCTCTTATCCCGACTA 1132  
Db 1404 CCTTTGGAGGTGAGAAAGTGGAAACAAATGTTTATTAACATCATTTCTTTGCTCGGA 1463  
Qy 1133 TGGTGTGCTGCACTGCTCTTCTCATCAATTTCTAGCCATTTATACCAGTCTTCAAGAG 1192  
Db 1464 TTGTTATTTGCTGACTTCTTTATTAATGAATCTGATCTCTGGGAGAGGATCTTCAGAG 1523  
Qy 1193 CAATTCCTTTTGGAAACAATGTTGGCGGTTGTTGCAATCTGTTTATTTGTTATCTTCCTC 1252  
Db 1524 CTATTCCTTTTGGACACTGGTTGCCATATTTGGCCCTTTGTTGCTGCAATCTGTGCTC 1583  
Qy 1253 TAAATCTTTGTTGATCAATACTTGGCGGAATCTGTGAGTCAAGTCAAGCCACTTCTGTC 1312  
Db 1584 TGACGTTTATTTGGTGCATACTTTGTTTAAAGAAATCCATGGAACAC---CCAGTTC 1640  
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Db 1641 GAACCAATCAGATTCACGTCAGATCTCTGAACAGTCTGTCTACAGAGCCCTTGCCTG 1700  
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Db 1701 GTATTATCATCGGAGGATTTTGGCCCTTTGGCTGCAATCTTTATACAACTTTTCTTCA 1760  
Qy 1433 TCAGCTCTTCTGGGATATAGATCTATTTATGTTCTATGCTCTGATGCTGCTGCTGG 1492  
Db 1761 TGAATAGTATTTGGTCAACACAGATGATATACATGTTGGCTTCTTATTTCTGGTGTTA 1820  
Qy 1493 TTATCTGTGCAATTTGCACTGTCTGTGCACTATTTGTGTCACATATTTCTTCTAAATG 1552  
Db 1821 TCATTTTGGTTATTAACCTGTTCTGAGCAACTATATCTTCTTGTCTATTTCCACCTAT 1880  
Qy 1553 CAGAGATTTACGGTGGCAATGACAAAGTTTCTCTCTGCTGATCAACTGCAATCTATG 1612  
Db 1881 CAGAGATTTATCATTTGGCAATGCGTTCTATCTCTACGAGTGGCTTTACTGCGATTTAT 1940  
Qy 1613 TTACATGATTTCTTTACTACTATTTTCAACAAAGATGATGCTTATTTCAA 1672  
Db 1941 TCTTAATCTATGCAATGACACTACTTCTTTCAAACTGCAAGTACGGGAAACAGCAAGCA 2000  
Qy 1673 CATCATTTTACTTTGGATATATGGCGGATTTTAGCACAGCTTTGGGATAATGTTGGAG 1732  
Db 2001 CAATCTGATCTTTGGTTATACCATGATATGTTTGTATCTTCTTCTTTTACAGGA 2060  
Qy 1733 CGATTGGTTTACATGGGAAACAAGTCTTTTGTCCGAAAAATCTATATTAATGGAATTTG 1792  
Db 2061 CAATTGGCTCTTTGATGCTTTTGGTTTGTGTTTACCAAAATATACAGTGTGGTGAAGTTG 2120  
Qy 1793 ACTAGAGA 1800  
Db 2121 ACTGAAGA 2128

RESULT 6

US-11-072-512-1699

; Sequence 1699, Application US/11072512  
; Publication No. US20060029945A1  
; GENERAL INFORMATION:  
; APPLICANT: ISOGAI, TAKAO  
; APPLICANT: SUGIYAMA, TOMOYASU  
; APPLICANT: OTSUKI, TETSUJI  
; APPLICANT: WAKAMATSU, AI  
; APPLICANT: SATO, HIROYUKI  
; APPLICANT: ISHII, SHIZUKO  
; APPLICANT: YAMAMOTO, JUN-ICHI  
; APPLICANT: ISONO, YUUKO  
; APPLICANT: HIO, YURI  
; APPLICANT: OTSUKA, KAORU  
; APPLICANT: NAGAI, KEIICHI  
; APPLICANT: IRIE, RYOTARO  
; APPLICANT: TAMECHIKA, ICHIRO  
; APPLICANT: SEKI, NAOHICO  
; APPLICANT: YOSHIKAWA, TSUTOMU  
; APPLICANT: OTSUKA, MOTOYUKI  
; APPLICANT: NAGAHARI, KENJI  
; APPLICANT: MASUHO, YASUHIKO  
; TITLE OF INVENTION: Novel full length cDNA  
; FILE REFERENCE: 084335-0191  
; CURRENT APPLICATION NUMBER: US/11/072,512  
; CURRENT FILING DATE: 2005-03-07  
; PRIOR APPLICATION NUMBER: US 60/350,978  
; PRIOR FILING DATE: 2002-01-25  
; PRIOR APPLICATION NUMBER: JP 2001-379298  
; PRIOR FILING DATE: 2001-11-05  
; NUMBER OF SEQ ID NOS: 4096  
; SOFTWARE: PatentIn Ver. 2.1  
; SEQ ID NO 1699  
; LENGTH: 1878  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
US-11-072-512-1699

Query Match 10.9%; Score 226; DB 9; Length 1878;

Best Local Similarity 51.8%; Pred. No. 2.5e-25;

Matches 595; Conservative 0; Mismatches 535; Indels 18; Gaps 3;

Qy 656 AATATCTTGATCCGCTCTTTTTCACATCGGATTCATGTTTCAATTTTCAACTCCT 715  
Db 693 ACTATATCTGGAGTCTATGCTCATACCCACATTCAGTGTGTAGCATTTGAATCCC 752  
Qy 716 TCATGATGGTGAATCTTCTGTTGGGCTTAGTTTCAATGATTTTATGAGAACATTAAGAA 775  
Db 753 TGGTCATTTGTTCTCTTCTTATCTGGAATGGTAGCTATGATTTATGTTACGGACCTGCACA 812  
Qy 776 AAGATTATGCTCGGTACAGTAAAGAGGAGAAATGGATGATGGATAGACCTTAGGAG 835  
Db 813 AAGATATTGCTAGATATATTAATCAGATGGACTCTACGGAAGATGCCAG-----G 860  
Qy 836 ATGAATATGATGAAACAGGTGCATGAGATGTATTTAGACCATCAAGTCAACCACTGA 895  
Db 861 AAGATTTGGCTGGAACATTTGTTCAATGATATATTTCCGCTCCCAAGAAAGGATGC 920  
Qy 896 TATTTTCTCTCTGATGTTTCTGAGTGCAGATTTTGTGTTGCTGTCTCTCATCGTTATTA 955  
Db 921 TGCTATCAGTCTTCTAGGATCCGGACACAGATTTTAAATATGACCTTTTGTGACTCTAT 980  
Qy 956 TTGTTGCAATGATAGAAATTTATATCTAGAGGGGATCAATGCTCAGTAC---AGCCA 1012  
Db 981 TTTTGGCTTGGCTGGGATTTTGTGACCTGCCACCGAGGAGCGCTGATGAGTGTGCTG 1040  
Qy 1013 TATTTGCTATGCTGCTACGCTCTCCAGTGAATGTTTATTTGGAGGAAGTCTGTATGCTA 1072  
Db 1041 TGGTCTGTGGGTGCTGCTGGGCACTCCCTGCGAGCTATGTTGCTGCCAGATTTCTATAAGT 1100  
Qy 1073 GACAAAGGAGGAGATGGATAAAGCAGATGTTTATTTGGGGCATTCCTTATCCAGCTA 1132  
Db 1101 CCTTTGGAGGTGAGAAAGTGGAAACAAATGTTTATTAACATCATTTCTTCTGCTCGGA 1160

1133 TGGTGTGGGACCTGCTTCTTCAATATTCATAGCCATTTATACCATGCTTCAAGAG 1192  
1161 TTGTATTTGCTGACTTCTTTATATGAATCTGATCTCTGGGAGAAAGGATCTTCAGCAG 1220  
1193 CCATTCCTTTTGGAAACATGTCGCTGTTGTCATCTGTTTGTGTTTATCTTCTCCTC 1252  
1221 CTATTCCTTTTGGACACTGTTGTCATATGTCGCTTGTGTTCTGCAATCTGTGCTC 1280  
1253 TAAATCTGTTGTPACAACTCTTGGCCGAATCTGTGAGTCAGCCCAACTTTCCTTGTG 1312  
1281 TGACGTTTATGTTGTCATCTTGTGTTTAAAGAAATGCAATGAACAC---CCAGTTC 1337  
1313 GTGTCAATGTCGCTGCTCTATACCGGAGAAATGTTTCAATGAGCCCTGCGGTTA 1372  
1338 GAACCAATCAGATTCACAGTCAGATTCCTGAACAGTCGTTCTACACGAAGCCCTTGCCTG 1397  
1373 TTGTTTCCCTGGGTGGAAATTTACCTTTTGTGTTCAATCTTTATTTGAATATGTTATCATCT 1432  
1398 GTATTATCATGGAGGATTTTGGCTTTGGCTGCACTTTTATACAACTTTTCTTCAATTC 1457  
1433 TCACGCTCTTTCTGGGCATATAAGATCTATTATGTCATGCTTCAATGATGCTGTGCTG 1492  
1458 TGAATAGTATTTGTCACACAGATGTTATACATGTTTGGCTTCTTATTTCTGCTGTTA 1517  
1493 TTATCTGTGCAATGTCAGTCTGTGTGACTATTGTGTCATATTTTCTACTAAATG 1552  
1518 TCATTTTGTGTTATTAACCTGTTCTGAAGCAACTATATCTTTTGTGCTATTTCCACCTATGTG 1577  
1553 CAGAAGATTAACCGTGGCAATGGAAGTTTCTCTGCTGTCATCAACTGCAATCTATG 1612  
1578 CAGAGGATTAATCATGGAATGGGCTTCATCTTACAGTGGCTTTACTGCAATTTAT 1637  
1613 TTTACATGATTTCTTTTACTACTATTTTTCAAAACAAAGATGATGCTTATTTTCAAA 1672  
1638 TCTTAATCTATGCAATGACTACTTCTTTTCAAACTGCAGATCAGCGGAACAGCAAGCG 1697  
1673 CATCATTTTACTTTGGATATATGCGGTATTTAGCAGAGCTTGGGATATGTTGTGGAG 1732  
1698 CAATTTCTGATCTTTGGTTATACCAATGATAATGTTTGTGATCTTCTTTTACAGGAA 1757  
1733 CGATTGTTTACATGGGAACAGTCCCTTTGTCGGAATAATCTATCTAATGTTGAAATG 1792  
1758 CAATGGCTCTTTGTATGCTTTTGGTTTGTGTACCAAAATATACAGTGTGTTGAGGTTG 1817  
1793 ACTAGAGA 1800  
1818 ACTGAAGA 1825

RESULT 7  
US-11-240-769-50  
; Sequence 50, Application US/11240769  
; Publication No. US20060036089A1  
; GENERAL INFORMATION:  
; APPLICANT: Soppet et al.  
; TITLE OF INVENTION: 33 Human Secreted Proteins  
; FILE REFERENCE: P2037Plc2  
; CURRENT APPLICATION NUMBER: US/11/240.769  
; CURRENT FILING DATE: 2005-10-03  
; PRIOR APPLICATION NUMBER: 09/997,131  
; PRIOR FILING DATE: 2001-11-30  
; PRIOR APPLICATION NUMBER: 09/628,508  
; PRIOR FILING DATE: 2000-07-28  
; PRIOR APPLICATION NUMBER: PCT/US00/03062  
; PRIOR FILING DATE: 2000-02-08  
; PRIOR APPLICATION NUMBER: 60/119,468  
; PRIOR FILING DATE: 1999-02-10  
; NUMBER OF SEQ ID NOS: 173  
; SOFTWARE: Patent In Ver. 2.0  
; SEQ ID NO 50  
; LENGTH: 1094  
; TYPE: DNA  
; ORGANISM: Homo sapiens

US-11-240-769-50  
Query Match 9.2%; Score 190.4; DB 9; Length 1094;  
Best Local Similarity 52.2%; Pred. No. 5.4e-20;  
Matches 494; Conservative 0; Mismatches 446; Indels 6; Gaps 3;  
QY 853 CAGGTGCATGGAGATGTTATTTAGACCATCAAGTCCACCACTGATATTTTCTCTCTGATT 912  
Db 9 CCGGTGCACGGGAGCGTCTTCAGGCCCCCCCAGTACCCCATGATCTCTAGCTCCCTGCTG 68  
QY 913 GGTTCCTGAGATGTCAGATATTTGCTGTGCTCTCATCGTTTATTTATTTGTCATGATGAA 972  
Db 69 GGTTCAGGATTCAGCTGTTCTGTATGATCTCTCATGTCATCTTTCTAGCCATGCTGG 128  
QY 973 GATTTATATCTAGAGA---GGGATCAATGCTCAGTACAGCCATATTTGTCATGCTGCT 1029  
Db 129 ATGCTGTCGCCCTCCAGCCGGGAGCTCTCATGACCACAGCCCTGCTTCTCTTCATGTTTC 188  
QY 1030 AGTCTCCAGTGAATGTTTATTTGGAGGAAGTCTGTATGCTAGACAAGGAGAAAGAGA 1089  
Db 189 ATGGGGGTGTTTGGCGGATTTCTGCTGCCGCTCTGTACGGCACTTTAAAGGCCATCGG 248  
QY 1090 TGGATAAAGCAGATGTTTATTTGGGCAATCTTATCCAGCTATGTTGTGTCATGCTGCC 1149  
Db 249 TGGAGAAAGGAGCCCTTCTGTACGGCAACTCTGTACCTGGTGTGTTTGGCATCTGC 308  
QY 1150 TTCTTCATCAATTTCAATGCCATTTATACCAATGCTTCAAGAGCCATTCCTTTTGGNACA 1209  
Db 309 TTGCTATTTGAATTTGCTTCAATTTGGGAAAGCACTCATCAGAGCGGTGCCCTTCCACC 368  
QY 1210 ATGTTGGCGCTTGTGTCATCTGTTTGTGTTTATTTCTCTCTAAATCTTGTGTTGATCA 1269  
Db 369 ATGTTGGCTCTGCTGTGCAATGTG---GTTCCGGAATCTCTGCGCCCTGCTACTTGGCTA 427  
QY 1270 ATATTTGGCGGAAATCTGTGAGTCCAGCCCAACTTTCTTGTGCTGTCATGTCGCTCT 1329  
Db 428 CTACTTGGCTTCCGAAAGCAG---CCATATGACAACCTGTGCGCAACCAACAGATTCCC 485  
QY 1330 CGTCTATACCGGAGAAATGTTTCATGGAGCCCTGCGTTATGTTTTCCTGCTGGGTGA 1389  
Db 486 CGGAGATCCCGGAGCAGCGGTGTACATGAAACCGATTTGTGGGCAATCTCATGCTGGG 545  
QY 1390 ATTTTACCTTTTGGTTCAATCTTTATTTGAAATGTAATTTTCACTTTTCACTGCTTTT 1449  
Db 546 ATCTTGGCTTCCGCGCCATGTTTCACTGAGCTCTTCTTCACTTTCAGTGTATCTGGAG 605  
QY 1450 TATAAGATCTATTATGTCATGCTTTCATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 1509  
Db 606 AATCAGTCTTATTAACCTTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 665  
QY 1510 ACTGCTGTGACTATTTGTGTCATATTTTCTACTAAATGCAAGATTAACCGGTGG 1569  
Db 666 TGTTCACAAATCAGCATCGTCAATGTTTCTTCCAGCTGTGTCAGAGGATTAACCGCTGG 725  
QY 1570 CAATGGAACAAGTTTCTCTGCTGCTCAATCACTGCAATCTATGTTTACATGATTTCTCTT 1629  
Db 726 TGTGGAAGAAATTTCTAGTCTCGGGGCTCTGCAATCTTACCTCTGTTTATGCAATC 785  
QY 1630 TACTACTATTTTTCAAAACAAAGATGATGCTTATTTTCAAAACATCAATTTTACTTTGGA 1689  
Db 786 TTTTATTTTCAACAAAGTGGACATCGTGGAGTTCATCCCTCTCTCTCTCTCTCTCTCTG 845  
QY 1690 TATATGCGGTATTTAGCAGAGCTTGGGATTAATGTTGGAGCGATGCTGTATCATGGA 1749  
Db 846 TACAGCGCCCTCATGCTTGTGCTTCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 905  
QY 1750 ACAAGTGCCTTTGTCGAAAAATCTATCTAATGTTGAAAAATGACT 1795  
Db 906 GCCTACATGTTTGTTCGCAAGATCTATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 951

RESULT 8  
US-11-240-769-49

```
; Sequence 49, Application US/11240769
; Publication No. US20060036089A1
; GENERAL INFORMATION:
; APPLICANT: Soppet et al.
; TITLE OF INVENTION: 33 Human Secreted Proteins
; FILE REFERENCE: P2037P1C2
; CURRENT APPLICATION NUMBER: US/11/240,769
; CURRENT FILING DATE: 2005-10-03
; PRIOR APPLICATION NUMBER: 09/997,131
; PRIOR FILING DATE: 2001-11-30
; PRIOR APPLICATION NUMBER: 09/628,508
; PRIOR FILING DATE: 2000-07-28
; PRIOR APPLICATION NUMBER: PCT/US00/03062
; PRIOR FILING DATE: 2000-02-08
; PRIOR APPLICATION NUMBER: 60/119,468
; PRIOR FILING DATE: 1999-02-10
; NUMBER OF SEQ ID NOS: 173
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 49
; LENGTH: 1821
; TYPE: DNA
; ORGANISM: Homo sapiens
; US-11-240-769-49

Query Match      8.3%; Score 172; DB 9; Length 1821;
Best Local Similarity 52.7%; Pred. No. 3.3e-17;
Matches 519; Conservative 0; Mismatches 445; Indels 20; Gaps 6;

QY 667 CCGTCCTTTTCAACATCGGATTCATTGGTTTCAATTTTCAATCTCTTCATGATGGT 726
DB 537 CTGACCATGAGTGCAGTCAGATCCACTGGTTTCTATCAATTAACCTCGTTGTTGGTGC 596
QY 727 ATCTCTTGGTGGGCTTAGTTTCATGATTTTAAAGCAATTAAGAAAGATTTATGCT 786
DB 597 TTCTTCTCTGTCAGGTATCTCGAGCATGATTAATCAATCGGACCCCTCGGAAGACATTC 656
QY 787 CGGTACAGTAAAGAGGAAGAAATGATGATGATGATGATGATGATGATGATGATGATG 846
DB 657 AACTACACACAGGAGATGACATTTGA-----AGACACCATGAGGAGTCTGGG 704
QY 847 TGGAAACAGGTGATGAGGAGATGATTTTA-GACCATCAAGTCAACCCACTGATATTTTCCTC 905
DB 705 TGGAAATGTTGTCAGCGGAGCGTCTTCAGGCGCCCTCCAGCATCCCATGATCCTCAGCTC 764
QY 906 TCTGATGTTGTTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 965
DB 765 CTTGCTGGGCTCAGGATTCAGCTGTTCTGTATGATGATGATGATGATGATGATGATGAT 824
QY 966 GATAGAGATTTATATAGTACAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 1022
DB 825 GCTTGGGATGCTGTCGCCCTCCAGCGGGAGCTCTCATGACCAAGCTGCTTCTTCTT 884
QY 1023 TGCTGCTACGTTCCAGTGAATGGTTATTTTGGAGGAAGTCTGTATGATGATGATGATG 1082
DB 885 CATGTTTATGCGGGGTGTTGGCGGATTTTCTGCTGCGGCTGTGTACCGCACTTTAAAGG 944
QY 1083 AAGGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1142
DB 945 CCATCGGTGGGAAGAAAGAGAGCGCTCTGTACGCGCAACTCTGTACCCCTGTTGGTTTGG 1004
QY 1143 CACTGCCCTTCTTCATCAATTTTCATAGCCATTTATTAACCATGCTTCAAGAGCCATTCCT 1202
DB 1005 CATCTGCTTGTATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1064
QY 1203 TGGAAACATGTTGGCGGTTTGTGATGATGATGATGATGATGATGATGATGATGATGAT 1262
DB 1065 TCCCAACCATGTTGGCTCTGCTGTGATGATG-CTTCGGGATCTCCCTGCGCTCTACT 1123
QY 1263 TGGTACATATCTTGGCCGAATCTGTGAGTCAAGCCCACTTCTTCTGCTGTTCAATGTC 1322
DB 1124 TGGGCTACTTCTCGGCTTCGGAAGAGCAG--CCATATGACAAACCCCTGTGCGCACCA 1181
QY 1323 TGTGCTGCTCTATATACCGGAGAAAAATGGTTTCATGAGCCTGCGGTTATTTGTTGCT 1382
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Db 1182 GATTCCTCCGCGAGATCCCGGAGCGGTGGTACATGAACCGATTTGTGGCATCCTCAT 1241
QY 1383 GGGTGGAAATTTTACCTTTTGGTTTCAATCTTTATTTGAAATGATTTTCACTTTTCACT 1442
Db 1242 GGTCTGGGATCTTGGC-TTCGGCGCCATGTTTCATCAGCTCTTCTTCACTTTCACTGCTAT 1300
QY 1443 CTGGGCATATAAGATCTATTATGTTCTATGCTTCTATGCTTCTATGCTGCTGCTGCTG 1502
Db 1301 CTGGGAGATCAGTTCTATTACCTCTTTGGTCTCTGTTCTGTTCTGTTCTGTTCTGTT 1360
QY 1503 CATTTGTAATCTGCTGTGTAATTTGTTGTCACATATTTTCTACTAAATGCAAGATTA 1562
Db 1361 GGTATCTCTGTTTCAAAATCAGCATCGTCAATGGTGTACTTCCAGCTGTGTGTCAGAGGATTA 1420
QY 1563 CCGTGGCAATGACAAATTTTCTCTCTGCTGTCATCACTGATCTATGTTTACATGTA 1622
Db 1421 CCGCTGGTGGGAGAAATTTCTTAGTCTCCGGGGCTCTGTCATTTCTACGTTCTGTTT 1480
QY 1623 TTCTCTTTTACTACTATTTTTCAT 1646
Db 1481 TGCATCTTTTATTTCTGTTAACA 1504

RESULT 9
US-11-240-769-21
; Sequence 21, Application US/11240769
; Publication No. US20060036089A1
; GENERAL INFORMATION:
; APPLICANT: Soppet et al.
; TITLE OF INVENTION: 33 Human Secreted Proteins
; FILE REFERENCE: P2037P1C2
; CURRENT APPLICATION NUMBER: US/11/240,769
; CURRENT FILING DATE: 2005-10-03
; PRIOR APPLICATION NUMBER: 09/997,131
; PRIOR FILING DATE: 2001-11-30
; PRIOR APPLICATION NUMBER: 09/628,508
; PRIOR FILING DATE: 2000-07-28
; PRIOR APPLICATION NUMBER: PCT/US00/03062
; PRIOR FILING DATE: 2000-02-08
; PRIOR APPLICATION NUMBER: 60/119,468
; PRIOR FILING DATE: 1999-02-10
; NUMBER OF SEQ ID NOS: 173
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 21
; LENGTH: 1816
; TYPE: DNA
; ORGANISM: Homo sapiens
; NAME/KEY: SITE
; LOCATION: (504)
; OTHER INFORMATION: n equals a,t,g, or c
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (1405)
; OTHER INFORMATION: n equals a,t,g, or c
; US-11-240-769-21

Query Match      8.2%; Score 169.4; DB 9; Length 1816;
Best Local Similarity 52.0%; Pred. No. 8e-17;
Matches 514; Conservative 7; Mismatches 448; Indels 20; Gaps 6;

QY 663 TGATCCGCTCTTTTTCACATCGGATTCATTGGTTTCAATTTTCAATCTCTTTCATGAT 722
DB 520 TTACTGACCATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 579
QY 723 GGTGATCTTCTTGGTGGGCTTAGTTTCAATGATTTTAAATGAGAACATTAAGAAAGATTA 782
DB 580 GGTCTTCTTCTGTCAGGTATCTTGAGCATGATTTATCATTCGGACCCCTCCGGAAGGACAT 639
QY 783 TGCTCGGTGATGATTAAGAGAGGAAGAAATGATGATGATGATGATGATGATGATGATGAT 842
DB 640 TGCCAACTACAAAGGAGGATGATGATTTGA-----AGACACCATGAGGAGTGC 687
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Db 630 CAATTTCATAGCCATTATTACCATGCTTCAAGAGCCATTCTTTTGGACAAATGGTG 573

RESULT 12

US-10-932-182A-478

; Sequence 478, Application US/10932182A

; Publication No. US20060046253A1

; GENERAL INFORMATION:

; APPLICANT: NAKAO, YOSHIHIRO

; APPLICANT: NAKAMURA, NORIHISA

; APPLICANT: KODAMA, YUKIKO

; APPLICANT: FUJIMURA, TOMOKO

; APPLICANT: ASHIKARI, TOSHIHIKO

; TITLE OF INVENTION: METHODS FOR ANALYZING GENES OF INDUSTRIAL YEASTS

; FILE REFERENCE: 030685-043

; CURRENT APPLICATION NUMBER: US/10/932,182A

; CURRENT FILING DATE: 2004-09-02

; NUMBER OF SEQ ID NOS: 197023

; SOFTWARE: PatentIn version 3.3

; SEQ ID NO 478

; LENGTH: 2019

; TYPE: DNA

; ORGANISM: Saccharomyces pastorianus

US-10-932-182A-478

Query Match 6.0%; Score 124.4; DB 7; Length 2019;

Best Local Similarity 46.4%; Pred. No. 4.7e-10;

Mismatches 567; Conservative 0; Mismatches 631; Indels 24; Gaps 4;

QY 577 GTTCCAAATACATAATCCAGATGTCATATTCAGTAAATGGAAGATGAGATGTAAC 636

Db 817 GATAAGATGATGAGCGTGTATTTTACCTACTCCGTCAATTCATGCTTGTGATACAGTT 876

QY 637 TTTGAAGATGATTTGACAAATATCTTGATCGCTCTTTTTCACATCGGATTCATTTGG 696

Db 877 TGGGCTACAGATGGGACAAATGATCTACAT-----ATTATGATCCGCAAAATTCATGG 930

QY 697 TTTTCAATTTTCACTCTTCATGATGTTGATCTCTTCTGGGCTAGTTTCAATGATT 756

Db 931 TTTTCTTTAAATTAATTTCTCCATCATCATCATTTTACATCATCTGTTGATTCATCTCT 990

QY 757 TTAATGAGAACATTAAGAAAGATTTATGCTCGGTACAGTAAAGAGGAAAGATGATGAT 816

Db 991 ATACTTGGGCTGTGAGAGTGAATTTGCGGTTATACGAA-----CTTCAC 1038

QY 817 ATGATAGAGACCTAGGAGATGAATATGATGGAACAGGTGTCATGAGATGATTTAGA 876

Db 1039 CTAGATAATGAATTTTCATGAGGATATCTGTTGGAATTTAGGTTCATGTTGATTTAGA 1098

QY 877 CCATCAAGTCCCACTGATATTTTCTCTCTGATTTGTTGTTGTTGTTGTTGTTGTTGTT 936

Db 1099 ACATCATCTAAATCAATGTTGTTATCTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTT 1158

QY 937 GTGCTCTCT---CATCGTTATTATGTTGCAATGATAGAGATTTATATATCTAGAGGGGA 993

Db 1159 ATGATCATATGATGATTTTCTTCTGCTGATTTAGGCTGTTATCCAGCTTCAAGAGGC 1218

QY 994 TCAATGCTCAGTACAGCCATTTTGTCTATGCTGCTACGTTCTCCAGTGAATTTTATTTT 1053

Db 1219 TCTTTTGCCAAACCGTATGTTGTTCTTTACGATTAATTCGGCTTGTAGTTCTTATACT 1278

QY 1054 GGAGGAGTCTGTATGCTAGACAGGAGGAGGATGATGATTAAGCAGATGTTTATTTGGG 1113

Db 1279 TCCATGGGTGCTATTAAGTTTTCATGGACCTTTATGGAAGGCTTAATTTGATTAATTA 1338

QY 1114 GCAATTCCTTATCCAGCTATGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTTGTT 1173

Db 1339 CCATCTCTGCTCTCTGCGGCAATTTTCTATTAATAGTAGCAATGAACTTCTTCTTATA 1398

QY 1174 TATTACCATGCTTCAAGAGCCATTCCTTTTGAACAATGGTGGCGGCTTTGTTGATCTGT 1233

Db 1399 TCTGGCAGTCTCTGGGTGTGATCCGACGAGAACGCTGTTTTCATATATCTTCTCTATGG 1458

QY 1234 TTTTGTGTATCTCTCTCTAAATCTTGTGGTGAATATCTTGGCCGAAATCTGTCAAGT 1293

Db 1459 TTTTGGGTGTCTATTTCATTTGCTGTTTGTGTTTGTGTTTGTGTTTGTGTTTGTGTTT 1518

QY 1294 CAGCCCAACTTTTCTCTGTCGTCGTCGTCGTCGTCGTCGTCGTCGTCGTCGTCGTCGTC 1353

Db 1519 TGGGATGACATCCAAACAAACAAACAAACAAACAAACAAACAAACAAACAAACAAACAA 1578

QY 1354 TTCAATGAGGCTCGGGTTATTTGTTTGGCTGGGTGGAATTTTACCTTTTGGTTCAATCTTT 1413

Db 1579 TACTTAAGGCAACACACAGGCAACCTTTATTGTCAGGCAATTTTCTCTTTTGGTTCAATAGCC 1638

QY 1414 ATTCAGATGATTTTCACTCTTCAGCTCTTCTGGGCATATAAGATCTATATGTCATGCGC 1473

Db 1639 GTTGAATTAATTTTCAATTTTCACTCTGATTTTGGTTTCAACAGATTTTGTATGTTGGT 1698

QY 1474 TTCAATGATGCTGGTGTGTTTATCTCTGTCATTTGTCGTCGTCGTCGTCGTCGTCGTCGTC 1533

Db 1699 TTTTCTACTCTTTTTCATTTTGTGTCGCTTTTAAACACCTCATTTGTTAACTATTTTTCATT 1758

QY 1534 ACATATTTTCTATAAATGACAGAAAGATTCGGTGGCAATGGAACAAAGTTTCTCTCTGCT 1593

Db 1759 ACGTATCATTTATGTTTGGAAATTTGGTTGGCAATTTGTTGGCAGTGGAGGATTTTATTATTGGT 1818

QY 1594 GCATCAACTGCAATCTATGTTTACATGTTTCTTTTACTTACTTATTTTTCACAAACAAG 1653

Db 1819 GGTT---TGGGATGTTTCAATTTATGTTTATGTTTCAATGCAATTTATTTTACAAATTCANA 1875

QY 1654 ATGATGCTGTTTATTTCAACATCATTTTACTTTGATATATATGCGGTATTTAGCACAGCC 1713

Db 1876 CTCGGTGGATTCGTCACAAATCATCTGATGTTGGATATTTCAATTTATGATATCTGTGCTA 1935

QY 1714 TTGGGATTAATGTTGGAGGATTTGTTTACATGGAACAAAGTTCCTTTGTCGGAATATC 1773

Db 1936 TGTTCGCTTGTACAGGGGCGATCGGCTTTTTCAGTAGCATGATATTTCAATTTAGAAAGAT 1995

QY 1774 TATACTAATGTGAAATTTGACT 1795

Db 1996 TATCTAGATTAAGTTCGAGT 2017

RESULT 13

US-10-932-182A-478

; Sequence 478, Application US/10932182A

; Publication No. US20060046253A1

; GENERAL INFORMATION:

; APPLICANT: NAKAO, YOSHIHIRO

; APPLICANT: NAKAMURA, NORIHISA

; APPLICANT: KODAMA, YUKIKO

; APPLICANT: FUJIMURA, TOMOKO

; APPLICANT: ASHIKARI, TOSHIHIKO

; TITLE OF INVENTION: METHODS FOR ANALYZING GENES OF INDUSTRIAL YEASTS

; FILE REFERENCE: 030685-043

; CURRENT APPLICATION NUMBER: US/10/932,182A

; CURRENT FILING DATE: 2004-09-02

; NUMBER OF SEQ ID NOS: 197023

; SOFTWARE: PatentIn version 3.3

; SEQ ID NO 478

; LENGTH: 2019

; TYPE: DNA

; ORGANISM: Saccharomyces pastorianus

US-10-932-182A-478

Query Match 6.0%; Score 124.4; DB 7; Length 2019;

Best Local Similarity 46.4%; Pred. No. 4.7e-10;

Mismatches 567; Conservative 0; Mismatches 631; Indels 24; Gaps 4;

QY 577 GTTCCAAATACATAATCCAGATGTCATATTCAGTAAATGGAAGATGAGATGTAAC 636

Db 817 GATAAGATGATGAGCGTGTATTTTACCTACTCCGTCAATTCATGCTTGTGATACAGTT 876

QY 637 TTTGAAGATGATTTGACAAATATCTTGATCGCTCTTTTTCACATCGGATTCATTTGG 696

Db 877 TGGCTACAGATGGGCAAGTATCTACAT-----ATTTAATCCGCAAAATTCATAGG 930  
Qy 697 TTTTCAATTTTCAACTCTCTCATGATGTGATCTTCTTGGTGGGCTAGTTTCAATGATT 756  
Db 931 TTTTCTTTAATTAATTTCTCATCATCATCATTTTACTATCATCTGTGGTTATTCATCT 990  
Qy 757 TTAATGAGAACATTAAGAAAGATTTATGCTCGGTACAGTAAAGAGGAAAGATGATAT 816  
Db 991 ATACTTCGGGTGTGAAGAGTGAATTTTGGCCGTATAACGAA-----CTTCAC 1038  
Qy 817 ATGATAGAGACCTAGGAGATGATATGAGTGAAGGAAACAGGTGTCATGAGATGATTTAGA 876  
Db 1039 CTAGATATGAATTTATGAGGATGACTGTTTGGAAATTAGTCAATGATGATTTAGA 1098  
Qy 877 CCATCAAGTACCCCACTGATATTTCTCTCTGATGGTTCTGATGTCAGATATTTGCT 936  
Db 1099 ACATCATTAATCAATGGTGTATCTGTCTGTGGGTTTCAGGTATTCATATTTCTG 1158  
Qy 937 GTGTCTCT---CATCGTTATTTATGTTGCAATGATAGAGATTTATATCTAGAGGGGA 993  
Db 1159 ATGATCATATGTAGCAATTTCTTGTCTGATTTAGGCTCGTATCACCAGCTTCAAGAGC 1218  
Qy 994 TCAATGCTCAGTACAGCAATTTGTCTATGCTGCTACGCTCCAGTGAATGGTTATTTT 1053  
Db 1219 TCTTTGCCAACCGTGTATTTGTTCTTTACGCATATTTCCGCTTTGAGGTTCTTACT 1278  
Qy 1054 GGAGGAAGTCTGTATGTAGCAAGGAGGAGATGATGATTAAGCAGATGTTTATTTGG 1113  
Db 1279 TCCATGGGTGTATAGTTTTCATGAGCTTATTTGGAAGCTAATTTGATTAATTA 1338  
Qy 1114 GCATTCCTTATCCAGTATAGGTGTGGCACTGCTTCTTCAATCAATTTCAAGCCATT 1173  
Db 1339 CCAATCCTGCTCTCGGCGCAATTTTCTATTAATAGTAGCAATGAATCTTCTTTATTA 1398  
Qy 1174 TATTAACATGCTTCAAGGCCATTTCTTTTGGACAATGTTGGCGGTTTGTGATCTGT 1233  
Db 1399 TCTGCGAGTCTCGGCTGTATCCAGCAAGAACGCTGTTTTCAATCTTCTCTTAGG 1458  
Qy 1234 TTTTGTGTATTTCTTCTCTAAATCTTGTGTGATCAATATCTGCGCGAAATCTGACAGT 1293  
Db 1459 TTTTGGGTGTCTATTCATTTCTGCTGTTCAATATATGCTACAAAAGTGAAC 1518  
Qy 1294 CAGCCCAACTTTCTGTGTGTCAATGCTGTGCTGTCTATACCGGAGAAAGATGG 1353  
Db 1519 TGGGATGAACATCCAAACAAAACCAACAGATGCTGCAAAAGTCCATTTTCAGCCTTG 1578  
Qy 1354 TTTCAATGGGCTGCGGTTATTTGTTGCTGCTGGAATTTTACCTTTTGTGTTCAATCTTT 1413  
Db 1579 TACTTAAGGCAACACAGGCAACCTTATTCAGGCAATTTCTCTTTGTTTCAATAGCC 1638  
Qy 1414 ATTGAATGTATTTTCACTCTTCACTCTTCTGCGCATATAGATCTATTTATGCTATGGC 1473  
Db 1639 GTTGAATTAATTTTCAATTTACTCCAGTTTATGTTTCAACAGATTTTGTATGTTGGT 1698  
Qy 1474 TTTCAATGATGCTGTGCTGTTATCTGTGCTGATGCTGCTGCTGCTGCTGCTGCTGCTG 1533  
Db 1699 TTTTCTACTCTTTTCAATTTTCTGCTGCTTAAACACCTCAATTTGTTAACTATTTTCA 1758  
Qy 1534 ACATATTTTCTACTAAATGCAAGATATTCAGGTTGCAATGCAAGTTTCTCTCTGCT 1593  
Db 1759 ACGTATCAATCATTTATGTTTGGAAATTTGGTTGGGAGTGGAGGATTTTATTTATTTGGT 1818  
Qy 1594 GCATCAACTGCAATCTATGTTTACATGATTTCTTTTACTACTATTTTCTTCAAAACAAAG 1653  
Db 1819 GGTT---TGGATGTTTCAATTTATGTTTATGTTTCAATGATTTTATTAACAAATTCAA 1875  
Qy 1654 ATGATGCTTATTTTCAAAACATTTTACTTTGATATATGCGGATTTTATGACAGCC 1713  
Db 1876 CTCGGTGGATTCGTACAAATCATATCTGATGTTGGATTTTCAATTTATGATATCTGTGCTA 1935  
Qy 1714 TTGGGATATGTTGGAGCAATTTGTTTACATGGGAACAGTGGCTTTGTCGGAATTC 1773  
Db 1936 TGTTCGCTGTGTCACAGGGGCGATCGGCTTTTTCAGTAGCATGATTTTCAATTAAGAGAT 1995

Qy 1774 TATACTAATGTGAAATTCAGT 1795  
Db 1996 TATTTCTAGAGTTAAAGTCGAGT 2017

## RESULT 14

US-10-932-182A-4690  
; Sequence 4690, Application US/10932182A  
; Publication No. US20060046253A1  
; GENERAL INFORMATION:  
; APPLICANT: NAKAO, YOSHIHIRO  
; APPLICANT: NAKAMURA, NORIHIISA  
; APPLICANT: KODAMA, YUKIO  
; APPLICANT: FUJIMURA, TOMOHIKO  
; APPLICANT: ASHIKARI, TOSHIO  
; TITLE OF INVENTION: METHODS FOR ANALYZING GENES OF INDUSTRIAL YEASTS  
; FILE REFERENCE: 030685-043  
; CURRENT APPLICATION NUMBER: US/10/932,182A  
; CURRENT FILING DATE: 2004-09-02  
; NUMBER OF SEQ ID NOS: 197023  
; SOFTWARE: PatentIn version 3.3  
; SEQ ID NO 4690  
; LENGTH: 2025  
; TYPE: DNA  
; ORGANISM: Saccharomyces pastorianus  
; US-10-932-182A-4690

Query Match 5.8%; Score 120; DB 7; Length 2025;  
Best Local Similarity 46.2%; Pred. No. 2.1e-09;  
Matches 562; Conservative 0; Mismatches 630; Indels 24; Gaps 4;

Qy 584 ATACTAAATCCAGATGTCTATATTCAGTAAATGGAAGAGTCAAGTGAATTTGAAG 643  
Db 830 ATAATCAAGTTTATTTTACCTATTCGTTAAATTCAGGAATCCCCACCTTCATGGGCTA 889  
Qy 644 ATCGATTTGCAATATCTTGATCCGCTCTTTTCAATCGGATTCATGTTTCA 703  
Db 890 CCAGATGGGCAATATTTTACA-----CGTTATGATCTCTTCCAAATGTTCTCT 943  
Qy 704 TTTTCAACTCTTCATGATGGTGTATCTTCTGTTGGGCTTAGTTTCAATGATTTTATGA 763  
Db 944 TAATTAATCTCTCTTGGTGTGCTATATCATCCGTCGTTATCCACTCTACTGTC 1003  
Qy 764 GAACATTAAGAAAGATTTATGCTCGGTACAGTAAAGAGGAGAAATGATGATGATA 823  
Db 1004 GCGCTTTGAAAAGTGTATTCGCTCGGTACAAACAGATTAACCTAGACGATGACTTCCA-- 1061  
Qy 824 GAGACCTAGGAGATGATATGATGGAACAGGTGATGAGATGATTTAGACCACATCAA 883  
Db 1062 -----AGAAGATTCAGGCTGGAATTAACCAACGATGATTTTCGTTCCGTTCCACAA 1111  
Qy 884 GTCACCACTGATATTTTCTCTCTGATTTGTTCTGATGTCAGATATTTGCTGTGTC 943  
Db 1112 GCCAGTCACTTAATGCTCTCCATTTTGGTTGTTTCAAGTGTTCATATTTTGTATGTC 1171  
Qy 944 TCATGCTATATTTGTTGCAATGATGATGATTTATATCTAGAGGGGATCAATGCTCA 1003  
Db 1172 CTTGTAGTATTTTGTGCTGCAATAGTTTCTCATCCCTAGCTCTAGAGGCTCGTTAG 1231  
Qy 1004 GTACGCCATTTTGTCTATGCTGCTACGTCCTCAGTGAATGTTTATTTGGAGGAA--- 1060  
Db 1232 CCACGTTATGTTTCAATTTATGCTTATTTGGATTTTGTGTTCTTACATCCATGG 1291  
Qy 1061 GTCTGTATGCTAGCAAGGAGGAGATGGAATAAAGCAGATGTTTATTTGGGCAATTC 1120  
Db 1292 GTATCTACAAATTTTTCGATGGTCCATCTAGGAGGCAATGATCATGACCCGCTTT 1351  
Qy 1121 TTAATCCAGCTATGTTGTGTCACCTCTTCTTCAATTTCAATGACCATTTATTTACC 1180  
Db 1352 TAGTTCCTGGAGCTATTTCTACTAGTAATCATTTGCACTGAACTTTTCTTAATGTTTCT 1411  
Qy 1181 ATGCTTCAAGAGCATTCCTTTTGGAAACATGTTGTCGCGTTTGTGTCATCTGTTTTTGT 1240

Db 1412 ATCTCTGGGTATTCAGCAAGTACTTTGTTTTTATGATATCTATGTTTTTAT 1471  
Qy 1241 TTAATCTTCTCTAAATCTTGTGGTACAAATCTTGGCGGAATCTGTGAGTCAAGCCCA 1300  
Db 1472 TCTCCATTCGGTTATCAATATGCTGTTCTGTTGTCAGGAAGAGATGTCACGGGATG 1531  
Qy 1301 ACTTTCCTTGTGCTGATGCTGTCGCTGCTGCTATACCGGAGAAAAAATGTTTCATGG 1360  
Db 1532 AGCATCCAAACGAAACGAAATTCGAAGCAAAATCCCTTCCAACTTGGTATCTGA 1591  
Qy 1361 AGCTCGCGGTATGTTGCTGCGGTGGAAATTTTACCTTTTGGTTCAATCTTATTGAAA 1420  
Db 1592 AAATCTACCGGCTACTTAAATGCTGTAATTTCCCTTCCGTTCTATGCGGTTGAAT 1651  
Qy 1421 TGTATTTCACTTCACGCTTTCTGGGATATAAGATCTATTTATGCTATGCGTTTCATGA 1480  
Db 1652 TGTATTTTATTTACCAAGTTTGTGTTCAATAAGATTTTCTACATGTTCCGGTTTCTAT 1711  
Qy 1481 TGCTGGTCTGTTATCTGTCATGTCGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 1540  
Db 1712 TTTTTCATTCCTTTTATTTGACGTTGACAACTCGTTGCTGCTGCTGCTGCTGCTGCTG 1771  
Qy 1541 TTCTACTAAATGCAAGATACCGGTGCGCAATGACAAAGTTTCTCTCTGCTGCTGCTGCA 1600  
Db 1772 ATTGTTATGCTAGAACTGGAAGTGGCAATGGAGGGGATTTATC---GTCCGAGGTG 1828  
Qy 1601 CTGCAATCTATGTTTACATGATTTCTTTTACTACTATTTTTCAAAAAAGATGATG 1660  
Db 1829 TCGGTTGCGCTGATGATTTATCCATTCATTTCACTAAATTTAAGTTAGGTG 1888  
Qy 1661 GCTTATTTCAACATCATTTTACCTTTGATATATGCGGATTTTATGACACGCTTGGGGA 1720  
Db 1889 GATTGCTTACCATTGTTTGTACTTGGGATATTTCTGTTATTTCAATGCTATGTTGTT 1948  
Qy 1721 TAAATGTTGGAGCGATTGGTTACATGGAACAAGTGCCTTTGTTCGAAAAATCTATATA 1780  
Db 1949 TAGTAACCTGGATCAATCGTTTCATTAGCAGCATGTTCTTTATAGAAAGATTTACTCAT 2008  
Qy 1781 ATGTGAAATTTGACTA 1796  
Db 2009 CCATCAAGTAGATTA 2024

## RESULT 15

US-10-932-182A-4690  
; Sequence 4690, Application US/10932182A  
; Publication No. US20060046253A1

GENERAL INFORMATION:  
; APPLICANT: NAKAO, YOSHIHIRO

; APPLICANT: NAKAMURA, NORIHISA  
; APPLICANT: KODAMA, YUKIKO

; APPLICANT: FUJIMURA, TOMOKO  
; APPLICANT: ASHIKARI, TOSHIHIKO

; TITLE OF INVENTION: METHODS FOR ANALYZING GENES OF INDUSTRIAL YEASTS  
; FILE REFERENCE: 030685-043

; CURRENT APPLICATION NUMBER: US/10/932,182A  
; CURRENT FILING DATE: 2004-09-02

; NUMBER OF SEQ ID NOS: 197023  
; SOFTWARE: PatentIn version 3.3

; SEQ ID NO 4690  
; LENGTH: 2025

; TYPE: DNA  
; ORGANISM: Saccharomyces pastorianus

US-10-932-182A-4690

Query Match 5.8%; Score 120; DB 7; Length 2025;  
Best Local Similarity 46.2%; Pred. No. 2.1e-09;

Matches 562; Conservative 0; Mismatches 630; Indels 24; Gaps 4;

Qy 584 ATACTAAATCCAGATGTCATATTCAGTAAATGAAAAAGTCAGATGTGAAATTTGAAG 643

Db 830 ATAATGAAGTTATTTTACCTATTGGTTAAATTCGAGGAATCCCCCACTTCATGGGCTA 889

Qy 644 ATGCAATTTGAACAATATCTTTGATCCGCTCTTTTTCACATCGGATCAATTTGGTTTTCAA 703  
Db 890 CCAGTGGGCAAAATATTTACA-----CGTTATGATCTCTCTATCCAATGGTTCTCCT 943  
Qy 704 TTTTCAACTCTTCATGATGATCTTCTTGGTGGGCTTAGTTTCAATGATTTTAAATGA 763  
Db 944 TAAATTAATCTCTCTTTGGTGGTGGCTATATATCATCCGCTGTTATCCACTCACTATGTC 1003  
Qy 764 GAACATTTAAGAAAGATTTATGCTCGGTPACAGTAAAGAGGAAGAAATGGATATGGATA 823  
Db 1004 GCGCTTTGAAGAGTGAATTCGCTCGGTACCAACGAGTTTAAACCTAGACGATGACTTCCA-- 1061  
Qy 824 GAGACCTAGGAGATGAATATGAGTGAACAGGTCAGTGGAGATGATTTAGACCATCAA 883  
Db 1062 -----AGAAGATTTCAGGCTGGAAATTAACCAACGCGTATGTTTCCGTTTCCACAA 1111  
Qy 884 GTCAACCACATGATATTTTCTCTGATTCGTTCTGATGTCAGATATTTGCTGTGCTC 943  
Db 1112 GCCAGTCACTAATGCTCTCCATTTTGGTGGTTCAGSTGTTCAATATTTTGTATGATGTC 1171  
Qy 944 TCATCGTTATTTTGTGCAATGATAGAAGATTTATATCTAGAGGGGATCAATGCTCA 1003  
Db 1172 CTTGTAGTATTTTGTGCTGCAATAGTTTCTCTATCACCTAGCTCAGAGGCTCGTTAG 1231  
Qy 1004 GTACAGCATATTTTGTCTATGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTG 1060  
Db 1232 CCAGGTTATGTTCAATTTCTATATGCTTATTTGGATTTTGGTTCCTACACATCCATGG 1291  
Qy 1061 GTCTGTATGCTAGCAAGGAGGAGATGGAATAAGACAGATGTTTATTTGGGCAATTC 1120  
Db 1292 GTATCTACAATTTTTCGATGGTCCATCTGGAAGGCAATGATGATCATGACCCGCTTT 1351  
Qy 1121 TTATCCAGCTATGGTGTGTCACCTGCTCTTCATCAATTTTCATAGCCATTTATTAACC 1180  
Db 1352 TAGTTCTGAGGCTATTTCTACTAGTAATCATTCGACTGAACTTTTCTTAATGTTTCTCC 1411  
Qy 1181 ATGCTTCAGAGCATTCCTTTTGGACATGCTGGCCGTTTGTGCTGCTGCTGCTGCTGCTG 1240  
Db 1412 ATTCTTCTGCTGTTATTTCCAGCAAGTACTTTGTTTATTTGTTATTTGTTGTTTAT 1471  
Qy 1241 TTATTTCTCTCTAAATCTTTGTTGGTACATACTTGGCCGAAATCTGTGAGTCAAGCCCA 1300  
Db 1472 TCTCCATTCGTTATCAATTTGCTGTTCTCTGTTGCCAGGAGAGATGTCATGGGATG 1531  
Qy 1301 ACTTTCCTGTCGTCATGTCGTCCTCTATACCGGAGAAAAAATGTTTCATGG 1360  
Db 1532 AGCATCCAAACGAAACGAAATTTGCAAGACAAATCCCTTCCAACCTTGGTATCTGA 1591  
Qy 1361 AGCTCGGTTATTTGTTGCTGCTGGGATTTTACCTTTTGGTTCATCTTTATTTGAAA 1420  
Db 1592 AAACTCTACCGGCTACTTTAAATTTGCTGGTATTTTCCCTTCCGTTCTATTTGCGGTTGA 1651  
Qy 1421 TGTATTTCACTTTCACTCTTTCTGCGCATATAAGATCTATATGCTATGCTGCTTCATGA 1480  
Db 1652 TGTATTTTATTTACACAAGTTTGGTTCATTAAGATTTTCTACATGTTCCGGTTCTAT 1711  
Qy 1481 TGCTGTGCTGGTTATCTGTCGATTTGTCATGCTGCTGTCGTCATTTGTCGTCATGTCAT 1540  
Db 1712 TTTTTCATTCCTTTTATTTGACGCTGACCAACCTCGTTGGTCACTGTAATGATCACTTATC 1771  
Qy 1541 TTCTACTAAATGCAAGATACCGGTGGCAATGGAACAGTTTCTCTCTGCTGCTGCTCA 1600  
Db 1772 ATTGTTATGCTAGAACTGGAAGTGGCAATGGAGGGGATTTATC---GTCCGAGGTG 1828  
Qy 1601 CTGCAATCTATGTTTACATGATTTCTTTTACTACTATTTTTCAAAAAAGATGATG 1660  
Db 1829 TCGGTTGCGCTGATGATTTATCCATTCATTTCACTAAATTTAAGTTAGGTG 1888  
Qy 1661 GCTTATTTCAACATCATTTTACCTTTGATATATGCGGATTTTATGACACGCTTGGGGA 1720  
Db 1889 GATTGCTTACCATTGTTTGTACTTGGGATATTTCTGTTATTTCAATGCTATGTTGTT 1948









QY 254 AGTGTCTGGTATTAATTTAAAGATGATGTGATGCCAGCCACTTACTGTGAATTCAT 313  
DB |||||||  
QY 301 AGTGTCTGGATTAATTTAAAGATGATGTGATGCCAGCCACTTACTGTGAATTCAT 360  
DB |||||||  
QY 314 TTAGATAAAGAAAGAGAGATGATTTGTATATGCCATAAATAATCATTTACTGGTACCAG 373  
DB |||||||  
QY 361 TTAGATAAAGAAAGAGAGATGATTTGTATATGCCATAAATAATCATTTACTGGTACCAG 420  
DB |||||||  
QY 374 ATGTACATAGATGATTTACCAATATGGGGTATTTGTGTGAGGCTGATGAAATGGAGAA 433  
DB |||||||  
QY 421 ATGTACATAGATGATTTACCAATATGGGGTATTTGTGTGAGGCTGATGAAATGGAGAA 480  
DB |||||||  
QY 434 GATTACTATCTTGGACCTTAATAAATCTTGAATAGCTTTTAAATGGAATCGAATTCAT 493  
DB |||||||  
QY 481 GATTACTATCTTGGACCTTAATAAATCTTGAATAGCTTTTAAATGGAATCGAATTCAT 540  
DB |||||||  
QY 494 GATGTTAATCTAACTAGTGAAGGAAGGTGAATCTGGTTTCAAAATCTAATAATCCAGATG 553  
DB |||||||  
QY 541 GATGTTAATCTAACTAGTGAAGGAAGGTGAATCTGGTTTCAAAATCTAATAATCCAGATG 600  
DB |||||||  
QY 554 TCATATTCAGTAAATGAAAGATCAGATGTGAATTTGAAGATCGAATTTGACAAATAT 613  
DB |||||||  
QY 601 TCATATTCAGTAAATGAAAGATCAGATGTGAATTTGAAGATCGAATTTGACAAATAT 660  
DB |||||||  
QY 614 CTTGATCCGCTCTTTTCAACATCCGATTCATTTGGTTTCAATTTTCAATCTCTTCATG 673  
DB |||||||  
QY 661 CTTGATCCGCTCTTTTCAACATCCGATTCATTTGGTTTCAATTTTCAATCTCTTCATG 720  
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QY 674 ATGGTATCTTCTTGGTGGGCTTAGTTTCAATGATTTTAAATGAGAAATTAAGAAAGAT 733  
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QY 721 ATGGTATCTTCTTGGTGGGCTTAGTTTCAATGATTTTAAATGAGAAATTAAGAAAGAT 780  
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QY 734 TATGCTCGGTACAGTAAAGAGAAAGATGATGATATGATAGACCTTAGAGATGAA 793  
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QY 781 TATGCTCGGTACAGTAAAGAGAAAGATGATGATATGATAGACCTTAGAGATGAA 840  
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QY 794 TATGATGGAAACAGGTGCATGGAGATGATTTAGACCATCAAGTCAACCACTGATATTT 853  
DB |||||||  
QY 841 TATGATGGAAACAGGTGCATGGAGATGATTTAGACCATCAAGTCAACCACTGATATTT 900  
DB |||||||  
QY 854 TCCTCTCTGATGGTTCTGGATGTCAGATATTTGCTGTGCTCTCATCTGATTAATTTGTT 913  
DB |||||||  
QY 901 TCCTCTCTGATGGTTCTGGATGTCAGATATTTGCTGTGCTCTCATCTGATTAATTTGTT 960  
DB |||||||  
QY 914 GCAATGATAGAGATTTATATCTAGAGAGGATCAATGCTCAGTACAGCCATTTGTC 973  
DB |||||||  
QY 961 GCAATGATAGAGATTTATATCTAGAGAGGATCAATGCTCAGTACAGCCATTTGTC 1020  
DB |||||||  
QY 974 TATGCTGTAGTCTCCAGTGAATGGTTATTTTGGAGGAAGTCTGTATGCTAGACAGGA 1033  
DB |||||||  
QY 1021 TATGCTGTAGTCTCCAGTGAATGGTTATTTTGGAGGAAGTCTGTATGCTAGACAGGA 1080  
DB |||||||  
QY 1034 GGAAGGATGATGATAAAGCAGATGTTATTTGGGCAATTCCTTATCCCACTGATGCTGT 1093  
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QY 1081 GGAAGGATGATGATAAAGCAGATGTTATTTGGGCAATTCCTTATCCCACTGATGCTGT 1140  
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QY 1094 GGCATGCTCTTCAATCAATTTTACAGCAATTTATACCAATGCTTCAAGAGCCATTCCT 1153  
DB |||||||  
QY 1141 GGCATGCTCTTCAATCAATTTTACAGCAATTTATACCAATGCTTCAAGAGCCATTCCT 1200  
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QY 1154 TTTGGAACAATGGTGGCGTTTGTGCAATCTGTTTTTGTGTTTTTGTATTTCTCTTAATCTT 1213  
DB |||||||  
QY 1201 TTTGGAACAATGGTGGCGTTTGTGCAATCTGTTTTTGTGTTTTTGTATTTCTCTTAATCTT 1260  
DB |||||||  
QY 1214 GTTGTGTAATCTTGGCCGAAATCTGTGAGTCAAGCCAACTTCTTGTGCTGTCAT 1273  
DB |||||||  
QY 1261 GTTGTGTAATCTTGGCCGAAATCTGTGAGTCAAGCCAACTTCTTGTGCTGTCAT 1320  
DB |||||||  
QY 1274 GCTGTGCTCTGCTTATACCGAGAAATATGGTTTCAATGAGGCTGCGGTTATTTGTC 1333  
DB |||||||  
QY 1321 GCTGTGCTCTGCTTATACCGAGAAATATGGTTTCAATGAGGCTGCGGTTATTTGTC 1380  
DB |||||||  
QY 1334 CTGGTGGAAATTTTACCTTTTGGTTCAATCTTTATTTGAATGATTTTCACTTCACTGCT 1393  
DB |||||||

DB 1381 CTGGTGGAAATTTTACCTTTTGGTTCAATCTTTATTTGAAATGATTTCACTTCACTGCT 1440  
QY |||||||  
QY 1394 TTCTGGGCATATAAGATCTATTATGCTATGCTCTCATGATGCTGGTGTGCTGCTTATCTG 1453  
DB |||||||  
QY 1441 TTCTGGGCATATAAGATCTATTATGCTATGCTCTCATGATGCTGGTGTGCTGCTTATCTG 1500  
DB |||||||  
QY 1454 TGCATTTGATCTGCTGTGTGATCTATTGCTGTGACATATTTTCTACTAAATGCAAGAT 1513  
DB |||||||  
QY 1501 TGCATTTGATCTGCTGTGTGATCTATTGCTGTGACATATTTTCTACTAAATGCAAGAT 1560  
DB |||||||  
QY 1514 TACCGTGGCAATGGAACAAGTTTCTCTGCTGTCATCAACTGCAATCTATGTTTACATG 1573  
DB |||||||  
QY 1561 TACCGTGGCAATGGAACAAGTTTCTCTGCTGTCATCAACTGCAATCTATGTTTACATG 1620  
DB |||||||  
QY 1574 TATTCCTTTTACTACTATTTTCAAAACAAAGATGATGCTTATTTTCAAAACATCATTT 1633  
DB |||||||  
QY 1621 TATTCCTTTTACTACTATTTTCAAAACAAAGATGATGCTTATTTTCAAAACATCATTT 1680  
DB |||||||  
QY 1634 TACTTTGGATATATGCGGTTATTTAGCAGAGCTTTGGGATATGCTGAGCGATTTGGT 1693  
DB |||||||  
QY 1681 TACTTTGGATATATGCGGTTATTTAGCAGAGCTTTGGGATATGCTGAGCGATTTGGT 1740  
DB |||||||  
QY 1694 TACATGGGAACAAGTGCCTTTTGTCCGAAATCTATATCTAATGTAATAATTTGACTAGAGA 1753  
DB |||||||  
QY 1741 TACATGGGAACAAGTGCCTTTTGTCCGAAATCTATATCTAATGTAATAATTTGACTAGAGA 1800  
DB |||||||  
QY 1754 CCCAAGAAACCTGGAACTTTGGATCAATTTCTTTTTCATAGGGGTGGAATTTGCAAGC 1813  
DB |||||||  
QY 1801 CCCAAGAAACCTGGAACTTTGGATCAATTTCTTTTTCATAGGGGTGGAATTTGCAAGC 1860  
DB |||||||  
QY 1814 AAAAAACAAAC 1827  
DB |||||||  
QY 1861 AAAAAACAAAC 1874  
DB |||||||

RESULT 3  
US-09-621-976-18829  
; Sequence 18829, Application US/09621976  
; Patent No. 6639063  
; GENERAL INFORMATION:  
; APPLICANT: Dumas Milne Edwards, J.B.  
; APPLICANT: Jobert, S.  
; APPLICANT: Giordano, J.Y.  
; TITLE OF INVENTION: ESTs and Encoded Human Proteins.  
; FILE REFERENCE: GENSET.054PR2  
; CURRENT APPLICATION NUMBER: US/09/621,976  
; CURRENT FILING DATE: 2000-07-21  
; NUMBER OF SEQ ID NOS: 19335  
; SOFTWARE: Patent.pm  
; SEQ ID NO 18829  
; LENGTH: 444  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
US-09-621-976-18829

Query Match 24.3%; Score 444; DB 3; Length 444;  
Best Local Similarity 100.0%; Pred. No. 7.4e-107;  
Matches 444; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 503 CTAACCTAGTGAAGGAAGGTGAACCTGGTTCCAAATCTAATAATCCAGATGTCATATTTCA 562  
DB |||||||  
QY 563 GTAAATGGAAGGAAGTGAATGTAAGATGCGATTTGAAATATCTTTGCAATATCTTTGATCCG 622  
DB |||||||  
QY 61 GTAAATGGAAGGAAGTGAATGTAAGATGCGATTTGAAATATCTTTGCAATATCTTTGATCCG 120  
DB |||||||  
QY 623 TCCTTTTTCACATCGGATTCATGTTGTTTCAATTTTCACTCTTCACTGATGCTGATC 682  
DB |||||||  
QY 121 TCCTTTTTCACATCGGATTCATGTTGTTTCAATTTTCACTCTTCACTGATGCTGATC 180  
DB |||||||  
QY 683 TTCCTTGGTGGCTTAGTTTCAATGATTTTAAATGGAACATTTAAGAAAGATTTATGCTCG 742  
DB |||||||

Db 181 TTTCTGGTGGCTTAGTTTCAATGATTTTAAATGAGACATTAAGAAAAGATTATGCTCGG 240  
QY 743 TACAGTAAAGAGGAAGAAATGGATGATGATGATAGAGACCTAGGAGATGAATATGGATGG 802  
Db 241 TACAGTAAAGAGGAAGAAATGGATGATGATGATAGAGACCTAGGAGATGAATATGGATGG 300  
QY 803 AAACAGGTGATGAGATGATTTAGACCATCAAGTCACCCACTGATATTTCTCTCTG 862  
Db 301 AAACAGGTGATGAGATGATTTAGACCATCAAGTCACCCACTGATATTTCTCTCTG 360  
QY 863 ATTGGTTCTGATGTCAGATATTGCTGTCTCTCATCGTTATTATTGTTGCAATGATA 922  
Db 361 ATTGGTTCTGATGTCAGATATTGCTGTCTCTCATCGTTATTATTGTTGCAATGATA 420  
QY 923 GAAGATTATATATCTAGAGGGGA 946  
Db 421 GAAGATTATATATCTAGAGGGGA 444

RESULT 4

US-09-513-999C-3708  
; Sequence 3708, Application US/09513999C  
; Patent No. 6783961  
; GENERAL INFORMATION:  
; APPLICANT: Dumas Milne Edwards, J.B.  
; APPLICANT: Duclert, A.  
; APPLICANT: Giordano, J.Y.  
; TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.  
; Patent No. 6783961  
; FILE REFERENCE: 59 US2 REG  
; CURRENT APPLICATION NUMBER: US/09/513,999C  
; CURRENT FILING DATE: 2000-02-24  
; PRIOR FILING DATE: 1999-02-26  
; NUMBER OF SEQ ID NOS: 36681  
; SOFTWARE: Patent.pm  
; SEQ ID NO 3708  
; LENGTH: 440  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; FEATURE:  
; NAME/KEY: CDS  
; LOCATION: 180..440  
; FEATURE:  
; NAME/KEY: misc\_feature  
; LOCATION: 151  
; OTHER INFORMATION: m=a or c  
; FEATURE:  
; NAME/KEY: misc\_feature  
; LOCATION: 155  
; OTHER INFORMATION: s=g or c  
; FEATURE:  
; NAME/KEY: misc\_feature  
; LOCATION: 162  
; OTHER INFORMATION: k=g or t  
; FEATURE:  
; NAME/KEY: misc\_feature  
; LOCATION: 184  
; OTHER INFORMATION: n=a, g, c or t  
; FEATURE:  
; NAME/KEY: misc\_feature  
; LOCATION: 323  
; OTHER INFORMATION: w=a or t  
; FEATURE:  
; NAME/KEY: misc\_feature  
; LOCATION: 343  
; OTHER INFORMATION: n=a, g, c or t  
; FEATURE:  
; NAME/KEY: misc\_feature  
; LOCATION: 397  
; OTHER INFORMATION: m=a or c  
; FEATURE:

; NAME/KEY: misc\_feature  
; LOCATION: 400  
; OTHER INFORMATION: m=a or c  
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; NAME/KEY: UNSURE  
; LOCATION: 2  
; OTHER INFORMATION: Xaa=Lys or Met or Arg or Thr  
; FEATURE:  
; NAME/KEY: UNSURE  
; LOCATION: 55  
; OTHER INFORMATION: Xaa=Ala or Asp or Gly or Val  
; FEATURE:  
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; OTHER INFORMATION: Xaa=Ala or Asp  
; FEATURE:  
; NAME/KEY: UNSURE  
; LOCATION: 74  
; OTHER INFORMATION: Xaa=Lys or Thr  
US-09-513-999C-3708  
Query Match 21.0%; Score 383.4; DB 3; Length 440;  
Best Local Similarity 95.7%; Pred. No. 6.7e-91;  
Matches 420; Conservative 5; Mismatches 8; Indels 6; Gaps 3;  
QY 591 TTGAAGATCGATTGAGAAATATCTTGATCCGTCCTTTTTCACACATCGGATTCATTTG 650  
Db 2 TTGAAGATCGATTGAGAAATATCTTGATCCGTCCTTTTTCACACATCGGATTCATTTG 61  
QY 651 TTTCAATTTTCAACTCCCTTCATGATGGTGATCTTCTTGGTGGGCTTAGTTTCAATGATTT 710  
Db 62 TTTCAATTTTCAACTCCCTTCATGATGGTGATCTTCTTGGTGGGCTTAGTTTCAATGATTT 121  
QY 711 TAATGAGAACATTAAGAAAAG---ATTATGCTCGGTACAGTAAAGAGAGAAATGGAT 766  
Db 122 TAATGAGAACATTAAGAAAAGAAATTAATGCTCGGTACAGTAAAGAGAGAAATGGAT 181  
QY 767 GAT-ATGATAGAGACCTAGGAGATGATATGATGGAACAGAGTGCATGGAGATGATTT 825  
Db 182 GAGATGATAGAGACCTAGGAGATGATATGATGGAACAGAGTGCATGGAGATGATTT 241  
QY 826 TAGACCATCAAGTCACCCACTGATATTTTCTCTGATTTGGTCTTGGATGTCAGATATT 885  
Db 242 TAGACCATCAAGTCACCCACTGATATTTTCTCTGATTTGGTCTTGGATGTCAGATATT 301  
QY 886 TGCTGTGTCCTCATCGTTATTTATTGTCATGATAGAGATTTATATCTGAGAGGGG 945  
Db 302 TGCTGTGTCCTCATCGTTATTTATTGTCATGATAGAGATTTATATCTGAGAGGGG 361  
QY 946 ATCAATGCTCAGTACAGCATATTTGTCATGCTGCTACGTCCT-CCAGTGAATGTTATT 1004  
Db 362 ATCAATGCTCAGTACAGCATATTTGTCATGCTGCTGATGCTCTCCAGTGAATGTTATT 421  
QY 1005 TTGGAGGAAGTCTGTATGC 1023  
Db 422 TTGGAGGAAGTCTGTATGC 440

RESULT 5

US-09-270-767-679/c  
; Sequence 679, Application US/09270767  
; Patent No. 6703491  
; GENERAL INFORMATION:  
; APPLICANT: Homburger et al.  
; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster  
; FILE REFERENCE: File Reference: 7326-094  
; CURRENT APPLICATION NUMBER: US/09/270,767  
; CURRENT FILING DATE: 1999-03-17  
; NUMBER OF SEQ ID NOS: 62517  
; SOFTWARE: Patentin Ver. 2.0  
; SEQ ID NO 679  
; LENGTH: 771  
; TYPE: DNA

; ORGANISM: Drosophila melanogaster  
US-09-270-767-679

Query Match 20.2%; Score 369.8; DB 3; Length 771;  
Best Local Similarity 67.7%; Pred. NO. 3.3e-87;  
Matches 518; Conservative 0; Mismatches 247; Indels 0

[illegible]

## RESULT 6

```

US-09-270-767-15961/c
; Sequence 15961, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
; APPLICANT: Homburger et al.
; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
; FILE REFERENCE: File Reference: 7326-094
; CURRENT APPLICATION NUMBER: US/09/270,767
; CURRENT FILING DATE: 1999-03-17
; NUMBER OF SEQ ID NOS: 62517
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 15961

```

; LENGTH: 771  
 ; TYPE: DNA  
 ; ORGANISM: D  
 US-09-270-767-1

Query Match	20.2%	Score 369.8	DB 3	Length 771
Best Local Similarity	67.7%	Pred. No. 3.3e-87	Indels 0	Gaps 0
Matches 518	Conservative	0	Mismatches 247	
Qy	563	GTAAAAATGGAAAAAGTCAGATGTGAATTTTGAAGATCGAATTTGACAAATATCTTGATCCG	622	
Db	765	GTCAACTGGNAGCCACGACAGGTGGAGTTCAAGAATCGATTTCGACAAGTACCTGGATCCC	706	
Qy	623	TCCTTTTTCACAAATCGGAATTCATTCGGTTTTCAAATTTTCAATCTCTTCATGATGTTGATC	682	
Db	705	AACTTCTTCGACGACAGGATCCACTGGTTTCAGCATCTTCAACAGCTTCATGATGGTCACT	646	
Qy	683	TTCTCTGGTGGCTTAGTTTCAATGATTTTAAATCAGAAACATTTAGAAAGAAATTTATGCTCGG	742	
Db	645	TTCTCTGGTGGCTTGGTGTCTCATGATTTCTGATCGGAATCTCTGGCGAAGGATTTATGCTCGG	586	
Qy	743	TACAGTAAAGAGAGAAATGGATGATATGGATAGAGACCTTAGGAGATGAATATGGATGG	802	
Db	585	TACAGTAAAGACGAGGAATCGACGACATGGAGCGAGATCTTGGTGATGAATACGGCTGG	526	
Qy	803	AAACAGGTGATGGAGATGATATTTAGACCAATCAAGTCAACCACTGATATTTTCTCTCTCTG	862	
Db	525	NAGCAGGTGCATGGCGATGTCTTCCGTCTCTCCGCCCAACACACTGTCTTCTCTGGCGTTG	466	
Qy	863	ATTGGTCTTGGATGTCAGATATTTGCTGCTCTCATCGTTATATATGTTGCAATGATA	922	
Db	465	GTGGCGCTGGATACCAACTGATTTTCGGTGTATATCTGTGTGATCATGTTTCGCCATAGTT	406	
Qy	923	GAAGATTTATATACCTCAGAGGGGATCAATGCTCAGTACAGCCATATTTGTCTATGCTGCT	982	
Db	405	GGTGAATTTGTACACGGNACGGGCTCCATGCTGTCCACGGCTATATTTGTATGTCGCCGC	346	
Qy	983	ACGTCTCCAGTGAATGGTTATTTTGGAGGAGTCTGTATGCTAGACAGGAGNAGGAGA	1042	
Db	345	ACCTCACCATCAATGGATACATTTGGAGGATCGCTCTATGCTCCGCCCTGGGTGGACGCATG	286	
Qy	1043	TGGATTAAGCAGATGTTTATTTGGGCATTCCTTATCCACGATATGTTGTCGGCACGTCC	1102	
Db	285	TGGATCCGACAGATGCTGGTGTCCGCTTTTACAGTTCACGTGGGTGTGTGCGGCACGGCT	226	
Qy	1103	TTCTTCATCAATTTCATAGCCATTTATACCATGCTTCAAGAGCCATTCCTTTTGGAAACA	1162	
Db	225	TTTCTGATCAACTTCATTTGCCATTTGGATATCAAGCCCTCGAGAGCCATTCCTTCGGTACC	166	
Qy	1163	ATGGTGGCGGTTGTTTGGCATCTGTTTTTTTGTATTTCTTCTCTAAATCTTGTGTTGGTACA	1222	
Db	165	ATGGTGGCGGTACAGTGCATCTGCCCTGTTTGTCTATCTCTGCTGCTGCTGCTGGTGGGTACT	106	
Qy	1223	ATACTTGGCCGAATCTGTACAGTTCAGCCCAACTTTTCTTGTGCTGTGCTCAATGCTGCTCT	1282	
Db	105	GTGCTGGGCGCGCAATCTGGAGCGCCCAACCGGACTTTTCCATGCGCGGTCAACGGGTGCCA	46	
Qy	1283	CGTCTTATACCGGAGAAAAATGGTTTCATGGAGCCTGCGGTTATT	1327	
Db	45	CGACCCCTTCTCCGAAAGAGAGTGTATCATGGAGCCACTGATTAAT	1	

## RESULT 7

```

RESOLUT /
US-09-513-999C-3502
; Sequence 3502, Application US/09513999C
; Patent No. 6783961
; GENERAL INFORMATION:
; APPLICANT: Dumas Milne Edwards, J.B.
; APPLICANT: Duclert, A.
; APPLICANT: Giordano, J.Y.
; TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
; Patent No. 6783961.
; FILE REFERENCE: 59.US2.REG

```

```
; CURRENT APPLICATION NUMBER: US/09/513,999C
; CURRENT FILING DATE: 2000-02-24
; PRIOR APPLICATION NUMBER: US 60/122,487
; PRIOR FILING DATE: 1999-02-26
; NUMBER OF SEQ ID NOS: 36681
; SOFTWARE: Patent.pm
; SEQ ID NO 3502
; LENGTH: 433
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: CDS
; LOCATION: 100..432
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: 86
; OTHER INFORMATION: m=a or c
US-09-513-999C-3502

Query Match      20.0%; Score 364.8; DB 3; Length 433;
Best Local Similarity 98.9%; Pred. No. 5.2e-86;
Matches 366; Conservative 1; Mismatches 3; Indels 0; Gaps 0;

QY 77 CACACGATCAAGATAAGAGGAGTGTCTTATGATGAATACCTCTTGGGCCCTACCAT 136
DB 64 CAGCAGTATCAAGATAAGAGGAGTGTCTTATGATGAATACCTCTTGGGCCCTACCAT 123
QY 137 AATCGTCAAGAAACATATAAGTACTTTTCACTTCCATTCTGTGTGGGGTCAAAAAAAGT 196
DB 124 AATCGTCAAGAAACATATAAGTACTTTTCACTTCCATTCTGTGTGGGGTCAAAAAAAGT 183
QY 197 ATCAGTCATTACCATGAATCTGGGAGAGCACTTCAAGGGGTTCGAATGGATTTAGT 256
DB 184 ATCAGTCATTACCATGAATCTGGGAGAGCACTTCAAGGGGTTCGAATGGATTTAGT 243
QY 257 GGTCTGGATTAATAATTAAGATGATGATGCCAGCACTTACTGTGAAATTTGATTTA 316
DB 244 GGTCTGGATTAATAATTAAGATGATGATGCCAGCACTTACTGTGAAATTTGATTTA 303
QY 317 GATAAGAAAAGAGAGATGATTTGATATGCCATAAAAAATCATTACTGGTACCAAGATG 376
DB 304 GATAAGAAAAGAGAGATGATTTGATATGCCATAAAAAATCATTACTGGTACCAAGATG 363
QY 377 TACATAGATGATTTACCATATGCGGTATGTTGGTGGCTGATGAAATGGAGAGAT 436
DB 364 TACATAGATGATTTACCATATGCGGTATGTTGGTGGCTGATGAAATGGAGAGAT 423
QY 437 TACTATCTTT 446
DB 424 TACTATCTTT 433

RESULT 8
US-09-270-767-28434
; Sequence 28434, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
; APPLICANT: Homburger et al.
; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
; FILE REFERENCE: File Reference: 7326-094
; CURRENT APPLICATION NUMBER: US/09/270,767
; CURRENT FILING DATE: 1999-03-17
; NUMBER OF SEQ ID NOS: 62517
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 28434
; LENGTH: 571
; TYPE: DNA
; ORGANISM: Drosophila melanogaster
US-09-270-767-28434

Query Match      12.6%; Score 230.8; DB 3; Length 571;
Best Local Similarity 72.7%; Pred. No. 1.1e-50;
Matches 298; Conservative 0; Mismatches 112; Indels 0; Gaps 0;

QY 1347 TACCTTTTGGTTCATCTTTATGAAATGATTTTCACTTTCACGCTTTCTGGGCATATA 1406
DB 1 TGCCCTTTGGATCCATCTTCAATGAGATGATCTTCACTTCCACCTCTTCTGGGCGTACA 60
QY 1407 AGATCTATTATGCTATGCTTATGCTTCAATGATGCTGGTCTGCTGATCTCTGTGCACTG 1466
DB 61 AGATCTATTATGCTATGCTTCAATGATGCTGGTCTGCTGATCTCTGTGCACTG 120
QY 1467 TCTGTGCACTATTGTGTGCAATATTTTCTACTAAATGCAAGAGATTACCGGTGGCAAT 1526
DB 121 TGTGGTCACTATGCTGTGCACTTCTCTGCTGCACTTCTCTGCTAAATGCCGAGATTACCGATGGCAGT 180
QY 1527 GGACAAGTTTCTCTCTGCTGCACTCACTGCAATCTATGTTTACATGATGATTTCTTTACT 1586
DB 181 GGACGAGTTTCAATGCTGCGGCTCCACGTCGATTTACGTTGACGCTATTTCTTCTATT 240
QY 1587 ACTATTTTTCAAAACAAAGATGATGCTTATTTTCAAAACATCAATTTTACTTTGGATATA 1646
DB 241 ACTTCTTCTTAAAACCAAAATGTTGGTCTGTTCCAAACGGCTTCTTACTTTGGCTACA 300
QY 1647 TGGCGGATTTAGCACACGCTTGGGGAATGTTGTGGAGCGATTTGGTACATGGAACAA 1706
DB 301 TGGCACTCTTCAGCGGCGCTTGGGCATTTATCTGCGCACCGTCGGCTATGTGGGCACGA 360
QY 1707 GTGCCCTTTGTCGAAAATCTTACTATGTAATGTAATTTGACTAGAGAGCC 1756
DB 361 ATCTCTTTGTGCGCAAAATCTATTCCAAATGTGAAATAGACTTAAGAGAGCC 410

RESULT 9
US-09-270-767-12633
; Sequence 12633, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
; APPLICANT: Homburger et al.
; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
; FILE REFERENCE: File Reference: 7326-094
; CURRENT APPLICATION NUMBER: US/09/270,767
; CURRENT FILING DATE: 1999-03-17
; NUMBER OF SEQ ID NOS: 62517
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 12633
; LENGTH: 1151
; TYPE: DNA
; ORGANISM: Drosophila melanogaster
US-09-270-767-12633

Query Match      12.6%; Score 230.8; DB 3; Length 1151;
Best Local Similarity 72.7%; Pred. No. 1.5e-50;
Matches 298; Conservative 0; Mismatches 112; Indels 0; Gaps 0;

QY 1347 TACCTTTTGGTTCATCTTTATGAAATGATTTTCACTTTCACGCTTTCTGGGCATATA 1406
DB 1 TGCCCTTTGGATCCATCTTCAATGAGATGATCTTCACTTCCACCTCTTCTGGGCGTACA 60
QY 1407 AGATCTATTATGCTATGCTTATGCTTCAATGATGCTGGTCTGCTGATCTCTGTGCACTG 1466
DB 61 AGATCTATTATGCTATGCTTCAATGATGCTGGTCTGCTGATCTCTGTGCACTG 120
QY 1467 TCTGTGCACTATTGTGTGCAATATTTTCTACTAAATGCAAGAGATTACCGGTGGCAAT 1526
DB 121 TGTGGTCACTATGCTGTGCACTTCTCTGCTGCACTTCTCTGCTAAATGCCGAGATTACCGATGGCAGT 180
QY 1527 GGACAAGTTTCTCTCTGCTGCACTCACTGCAATCTATGTTTACATGATGATTTCTTTACT 1586
DB 181 GGACGAGTTTCAATGCTGCGGCTCCACGTCGATTTACGTTGACGCTATTTCTTCTATT 240
QY 1587 ACTATTTTTCAAAACAAAGATGATGCTTATTTTCAAAACATCAATTTTACTTTGGATATA 1646
DB 241 ACTTCTTCTTAAAACCAAAATGTTGGTCTGTTCCAAACGGCTTCTTACTTTGGCTACA 300
QY 1647 TGGCGGATTTAGCACACGCTTGGGGAATGTTGTGGAGCGATTTGGTACATGGAACAA 1706
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Db 301 TGGCACTCTTCAGCGCGCCCTTGGCCATTATCTCGGCACCGTGGCTATGTGGCAGCA 360  
Qy 1707 GTGCTTTGTCCGAAAATCTATCTAATGTGAAAATTCAGCTAGAGACCC 1756  
Db 361 ATCTTTGTGCGCAAAATCTTCCATGTGAAAATAGACTRAGAGCCC 410

RESULT 10  
US-09-949-016-3623  
; Sequence 3623, Application US/09949016  
; Patent No. 6812339  
; GENERAL INFORMATION:  
; APPLICANT: VENTER, J. Craig et al.  
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED  
; WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF  
; FILE REFERENCE: CLO01307  
; CURRENT APPLICATION NUMBER: US/09/949,016  
; CURRENT FILING DATE: 2000-04-14  
; PRIOR APPLICATION NUMBER: 60/241,755  
; PRIOR FILING DATE: 2000-10-20  
; PRIOR APPLICATION NUMBER: 60/237,768  
; PRIOR FILING DATE: 2000-10-03  
; PRIOR APPLICATION NUMBER: 60/231,498  
; PRIOR FILING DATE: 2000-09-08  
; NUMBER OF SEQ ID NOS: 207012  
; SOFTWARE: FastSeq for Windows Version 4.0  
; SEQ ID NO 3623  
; LENGTH: 2391  
; TYPE: DNA  
; ORGANISM: Human  
US-09-949-016-3623

Query Match . 12.5%; Score 227.6; DB 3; Length 2391;  
Best Local Similarity 51.9%; Pred. No. 1.5e-49;  
Matches 596; Conservative 0; Mismatches 534; Indels 18; Gaps 3;

Qy 609 AATATCTTGATCCGFCCTTTTTCACATCGGATTCATGTGTTTCAATTTCACTCCT 668  
Db 996 ACTATATTCGAGTCTATGCTCATACCCACATTCAGTGGTTTAGCAATTGAATCCC 1055  
Qy 669 TCATGATGGTATCTCTTGTGGCTAGTTTCATGATTTTAAATGAGACATTAAGAA 728  
Db 1056 TGGTCATTTGCTCTCTCTTATCTGGAATGGTAGCTATGATTATGTACGGACATGCACA 1115  
Qy 729 AAGATTATGCTCGGTACAGTAAGAGGAAAGAAATCGATCATATGATAGACCTTAGGAG 788  
Db 1116 AAGATTATGCTAGATATATATCAGATGGACTCTACGGAGATGCCAG-----G 1163  
Qy 789 ATGAATATGGATGGAAAACAGGTGCATGGAGATGATTTTAGACCATCAAGTCAACCCTGA 848  
Db 1164 AAGAAATTTGGCTGGAAACTTGTTCATGTGTATATATCCGTCCTCCAAGAAAAGGGATGC 1223  
Qy 849 TATTTCTCTGATTTGGTCTGGATGTCAGATATTTGCTGTGCTCTCATCGTTATTA 908  
Db 1224 TGCTATCAGTCTTCTAGGATCCGGACACAGATTTTAAATGACCTTTGTGACTCTAT 1283  
Qy 909 TTGTTGCAATGATAGAAGATTTATATCTATGAGAGGGGATCAATGCTCAGTAC---AGCCA 965  
Db 1284 TTTTCGCTGCTGGATTTTGTACCTGCCAACCGAGGAGCGTGTAGCGTGTCTG 1343  
Qy 966 TATTTGTCTAGCTGCTAGCTCCAGTGAATGGTTTATTTGGAGAAAGTCTGTATGCTA 1025  
Db 1344 TGGTCTGTGGGTGCTGTGGGCAACCCCTGCAGGCTATGTTGCTGCCAGATTTCTATAAGT 1403  
Qy 1026 GACAGGAGGAGGAGATGATGAAGACGATGTTTATTTGGGCAATTCCTATCCAGCTA 1085  
Db 1404 CCTTTGGAGGTGAGAAGTGGAAAACAAATGTTTATTAACATCAATTTCTTTGCTCTGGGA 1463  
Qy 1086 TGGTGTGTGGCACTGCTCTTCTTCATCAATTTTCATAGCCATTTTATTAACATGCTCAAGAG 1145  
Db 1464 TTGTAATTTGCTCACTTCTTTATTAATGAATCTGATCTCTCTGGGAGAGGATCTTCAGCAG 1523

Qy 1146 CCATTCCTTTTGAACAATGGTGGCCGTTTGTGTCATCTGTTTTTTTGTATTCTTCTC 1205  
Db 1524 CTATTCCTTTTGGGACACTGGTTGGCCATATGTGCCCTTTGGTTCTGCAATATCTGTGGCTC 1583  
Qy 1206 TAAATCTTGTGTGTAACAATACCTTGGCCGAATCTGTGAGTCAGCCCAACTTTTCTTGTGTC 1265  
Db 1584 TGACGTTTATTTGGTCATCTTTGGTTTAAAGAAAGATGCCATTGAACAC---CCAGTTC 1640  
Qy 1266 GTGTCAATGCTGTGCTGCTCTATACCGGAGAAAAAATGGTTTCATGGAGCCCTGCGGTTA 1325  
Db 1641 GAAACCAATCAGATTCCACGTCAGATTCTCTGAACAGTCGTTCTACAGAAAGCCCTTGCCGT 1700  
Qy 1326 TTGTTTGGCTGGGTGGAATTTTACCTTTTGGTTTCAATCTTTATTTGAATGTATTTTCACT 1385  
Db 1701 GTATTATATGGAGGAGATTTTCCCTTTGGCTGCACTTTTATACAACCTTTTCTTCAATTC 1760  
Qy 1386 TCACGCTTTTCTGGGCATATAAGATCTATTATGCTGTATGCTTTCATGATGCTGTGCTGG 1445  
Db 1761 TGAATAGTATTTTGGTCACACACAGATGATTAATCATGTTTGGCTTCTTATTTCTGGTGTTA 1820  
Qy 1446 TTATCTGTGCAATTTGACGTCTGTGTGACTATTTGTGTCACATATTTTCTACTAAATG 1505  
Db 1821 TCATTTTGGTTATTTACCTGTTCTGAAGCAACTATATCTTCTTGGCTATTTCCACCTATGTG 1880  
Qy 1506 CAGAAGATTACCGTGGCAATGACAAAGTTTCTCTCTGCTGCATCAACTGCAATCTATG 1565  
Db 1881 CAGAGGATATCATTTGGCAATGGCTTTCATCTCTAGGAGTGGCTTTTACTGCACTTAT 1940  
Qy 1566 TTTTCAATGATTCCTTTTACTACTATTTTTTCAAAAACAAAGATGTATGGCTTATTTTCAA 1625  
Db 1941 TCTTAATCTATGCAGTACACTACTTCTTTTCAAAAACCTGCAGATCACGGGAACAGCA 2000  
Qy 1626 CATCATTTTACTTTGATATATGCGGTATTTAGCACAGCCTTGGGATATATGTTGTCGAG 1685  
Db 2001 CAATTCGTACTTTGGTTATATACCATGATATGTTTGGTATCTTCTTTTTCACAGAA 2060  
Qy 1686 CGATTGTTTACATGGGAACAAAGTGCCTTTCTCGGAAAAATCTATPACTAATGTGAAAAATTG 1745  
Db 2061 CAATGGCTTCTTGGCATGCTTTTGGTTTGTACCAAAATATACAGTGTGTGAGGTTG 2120  
Qy 1746 ACTAGAGA 1753  
Db 2121 ACTGAGAGA 2128

RESULT 11  
US-08-959-004-6  
; Sequence 6, Application US/08959004  
; Patent No. 6197543  
; GENERAL INFORMATION:  
; APPLICANT: Hillman, Jennifer L.  
; APPLICANT: Yue, Henry  
; APPLICANT: Corley, Neil C.  
; APPLICANT: Lal, Preeti  
; APPLICANT: Shah, Purvi  
; APPLICANT: Kaser, Matthew  
; TITLE OF INVENTION: HUMAN VESICLE MEMBRANE PROTEIN-LIKE  
; TITLE OF INVENTION: PROTEINS  
; NUMBER OF SEQUENCES: 11  
; CORRESPONDENCE ADDRESS:  
; ADDRESSEE: Incyte Pharmaceuticals, Inc.  
; STREET: 3174 Porter Drive  
; CITY: Palo Alto  
; STATE: CA  
; COUNTRY: USA  
; ZIP: 94304  
; COMPUTER READABLE FORM:  
; MEDIUM TYPE: Diskette  
; COMPUTER: IBM Compatible  
; OPERATING SYSTEM: DOS  
; SOFTWARE: FastSeq for Windows Version 2.0  
; CURRENT APPLICATION DATA:  
; APPLICATION NUMBER: US/08/959,004

Db	1689	GAACCAATCAGATTCCACGTCAGATTCTTGAAACAGTCGTTCTACACGAAGCCCTTGCCCTG	1748
Qy	1326	TTGCTTTGCCCTGGGCGAATTTTACCTTTTGGTTCAATCTTTATTTGAAATGATTTTCAATCT	1385
Db	1749	GTATATATATGCGAGGAGATTTTCCCTTTGGCTGCAATCTTTATACAACTTTTCTTCATTC	1808
Qy	1386	TCACGCTCTTTCTCGGCATATATAGATCTATTTATGTCCTATGGCTTCATGATGCTGGTCTGG	1445
Db	1809	TGAATAGTATTTTGGTCAACACAGATGATTTACATGTTTGGCTTCTCTATTTCTCGTGTTTA	1868
Qy	1446	TTATCCCTGTCATTTGTGACTGTCCTGTGTGACTATTTGTGTGACATATATTTTCTACTAAATG	1505
Db	1869	TCAATTTGGTATTATACCTGTTCTGGAAGCAACTATATCTTCTTGTCTATTTCCACCTATGTG	1928
Qy	1506	CACAAGATTACCGGTGCGCAATGCAACAAGTTTCTCTCTGCTGCATCAACTGCAATCTATG	1565
Db	1929	CAGAGATTATCATTTGGCAATGGCGTTCAATCTTAGAGTGGCTTTACTTGCAGTTTATTT	1988
Qy	1566	TTTTACATGTAATCCCTTTTACTACTATTTTTTCAAAAACAAAGATGTATGGCTTATTTCAAA	1625
Db	1989	TCCTTAATCTATGCACTACTCTCTTTTCAAAAACAGATCACGGGAACAGCAAGCA	2048
Qy	1626	CATCATTTTACTTTTGATATATGGCGGTTATTTAGCACAGCCTTGGGGAATATGTGTGGAG	1685
Db	2049	CAATTCGTACTTTGGTTATACCATGATAATGGTTTGTGATCTTCTTTTTCACAGGAA	2108
Qy	1686	CGATGCTTACATGCGAACAAAGTGCCCTTTGTCCGAAAAATCTATCTAAATGTCAAAAATTG	1745
Db	2109	CAATGGCTCTTTTGATGCTTTTGGTTTGTACAAAATATACAGTGTGTGAAAGTTG	2168
Qy	1746	ACTAGAGA 1753	
Db	2169	ACTGAAGA 2176	

RESULT 12

US-10-104-047-1699

; Sequence 1699, Application US/10104047

; Patent No. 6943241

; GENERAL INFORMATION:

; APPLICANT: HELIX RESEARCH INSTITUTE

; TITLE OF INVENTION: No. 6943241el full length cDNA

; FILE REFERENCE: H1-A0105

; CURRENT APPLICATION NUMBER: US/10/104,047

; CURRENT FILING DATE: 2002-03-25

; PRIOR APPLICATION NUMBER:

; PRIOR FILING DATE:

; NUMBER OF SEQ ID NOS: 4096

; SOFTWARE: PatentIn Ver. 2.1

; SEQ ID NO 1699

; LENGTH: 1878

; TYPE: DNA

; ORGANISM: Homo sapiens

US-10-104-047-1699

Query Match 12.4%; Score 226; DB 3; Length 1878;

Best Local Similarity 51.8%; Pred. No. 3.5e-49;

Matches 595; Conservative 0; Mismatches 535; Indels 18; Gaps 3;

Qy	609	AATATCTTGATCCGTCCTTTTTTCAACATPCGGATTCATTTGGTTTCAATTTTCAACTCCCT	668
Db	693	ACTATATCTGGAGTCTATGCTCTCATACCCACATTCAGTGGTTTAGCAATATGAAATCCC	752
Qy	669	TCATGATGTGATCTTCTTGGTGGCTTAGTTTCATGATTTTATGAGAACATTAAGAA	728
Db	753	TGCTCATTTCTCTTCTTATCTGGAATGGTAGTATGATGATTTTACGGACACTGCACA	812
Qy	729	AGAGTTATCTCGGTACAGTAAGAGAGAAATGGATGATATGGATAGAGACCTAGGAG	788
Db	813	AGATATCTAGATATATATCAGTGGATCTTACGGAAGATGCCAG-----G	860
Qy	789	ATGAATATGGATGGAAACAGGTGTCATGGAGATGTATTTTAGACCATCAAGTCAACCACTGA	848



861	AGAAATTGGCTGGAACCTTGTCATGGTGAAATATTTCCGTCTCTCAAGAAAAGGGATGC	920
849	TATTTTCTCTCTGATTTGGTTCTGGATGTCAGATAATTTGCTGTCTCTCATCGTTATTA	908
921	TGCTATCAGTCTTCTAGGATCCGGACACAGATTTTAAATTATGACCTTTGTGACTCTAT	980
909	TTGTTGCATGATAGAGATTATATCTCAGAGGGGATCAATGCTCAGTAC--AGCCA	965
981	TTTTCGCTTGCCCTGGGATTTTGTGCACCTGCCAACGAGAGCGCTGATCAGCTGTGCTG	1040
966	TATTTGCTATGCTGCTACGCTCTCCAGTGAAATGGTTATTTTGGAGGAAGTCTGTATGCTA	1025
1041	TGCTCTGTGGGTGCTCTGGGACCCCTGCAGGCTATGTTGCTGCCAGATTCTATAAGT	1100
1026	GACAGGAGGAGGAGATGGAATAAGCAGATGTTTATTTGGGCAATTCCTTATCCACGCTA	1085
1101	CCTTTGGAGGTGGAAGTGGAAAACAAATGTTTATTAACATCATTTCTTTGTCTCTGGGA	1160
1086	TGCTGTGTGGCACTGCCCTCTTCATCAATTTCTATAGCCATTTATTACCATGCTTCAAGAG	1145
1161	TTGTATTGCTGACTCTTTTATATGATCTGATCCTCTGGGAGAGGATCTTCAGCAG	1220
1146	CAATTCCTTTTGAACAAATGGTGGCGGTTGTTGTCATCTGTTTTTTTGGTTATTTCTCTC	1205
1221	CTATTCTTTTGGGACACTGGTTGGCCATATGGCCCTTTGGTTCTGTCATATCTGTGCTC	1280
1206	TAAATCTGTGTGTCACAATACTTTGGCCGAATCTGTCAAGTCAGCCCAACTTCTCTTGTG	1265
1281	TGACGTTTATTTGGTGCATCTTTGGTTTTAAGAAGAAATGCCATTTGAACAC--CCAGTTC	1337
1266	GTGTCGAATGCTGTGCCCTCGTCTTATACCGGAGAAAAATGGTTTCATGGAGCCCTGCGGTTA	1325
1338	GAACCAATCAGATTCCACGTCAGATTCCTGAAAGTGCTTCTACAGAAAGCCCTTGGCTG	1397
1326	TTGTTTCCCTGGGTGGAAATTTTACCTTTTGGTTCAATCTTTTATTTGAATGTTTCACTCT	1385
1398	GTATTATCATGGGAGGGAATTTGGCCCTTTGGCTGCACTTTTATACAACTTTTCTTCAATC	1457
1386	TCACGCTTTTCTGGGCATATAAGATCTATTATGTCATATGCTTATGGCTTCATGATGCTGCTGG	1445
1458	TGAAATGATTGTTGGTCACACAGATGTTATACATGTTTGGCTTCTTATTTCTGGTGTTTA	1517
1446	TTATCCTGTGCATGTGACTGTCTGTGTGACTATTTGTGTGCAATATTTTCTACTAAATG	1505
1518	TCAATTTGGTTATTACCTGTTCTGAAGCAACTATCTTCTTGTCTATTTTCCACCTATGTG	1577
1506	CAGAAGATTACCGTGGCAATGGACAAGTTTCTCTCTGCTGCAATCAACTGCAATCTATG	1565
1578	CAGAGGATTCATTTGGCAATGGCGTTTCATCTCTACAGTGGCTTTACTTGCAGTTTATTT	1637
1566	TTTTACATGTTATCCTTTTACTACTATTTTTTCAAAACAAAGATGTATGGCTTATTTCAAA	1625
1638	TCTTAATCTATGCAGTACACTACTCTCTTTTCAAAACCTGCAGATCACGGGAACAGCAAGC	1697
1626	CATCATTTTACTTTGGATATATGCGGTATTTAGCACAGCCCTTGGGGATATGTGTGGAG	1685
1698	CAATTCGTACTTTGGTTATACCAATGATAATGGTTTGTGATCTTCTTTTTCACAGGAA	1757
1686	CGATTTGGTTACATGGGAACAAGTGCCCTTTGTCGAAAAAATCTATPACTAATGTGAAAAATTG	1745
1758	CAATTGGCTCTTTGTATGCTTTTGGTTTGTGTACCAAAATATACAGTGTGGTGAAGGTTG	1817
1746	ACTAGAGA	1753
1818	ACTGAAGA	1825

RESULT 13  
US-09-270-767-14715  
; Sequence 14715, Application US/09270767  
; Patent No. 6703491  
; GENERAL INFORMATION:

```

; APPLICANT: Homburger et al.
; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
; FILE REFERENCE: File Reference: 7326-094
; CURRENT APPLICATION NUMBER: US/09/270,767
; CURRENT FILING DATE: 1999-03-17
; NUMBER OF SEQ ID NOS: 62517
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 14715
; LENGTH: 995
; TYPE: DNA
; ORGANISM: Drosophila melanogaster
US-09-270-767-14715

```

Query Match	8.8%	Score 161.2;	DB 3;	Length 995;
Best Local Similarity	59.3%;	Pred. No. 3.1e-32;		
Matches 274;	Conservative 0;	Mismatches 188;	Indels 0;	Gaps 0;
QY	48	TGCCCCGGACCCGGGGGAGCAGCAAGAACACACACGTATCAAGATATAAGAGGAGTTGTCT	107	
DB	532	TGTCACCTCTCCCAAGGCAGATGAGCACAAATCACAAGTACAATGACCGGGAGGCTGGTAC	591	
QY	108	TATGGATGAATACTGTTGGCCCTTACCATAATCGTCAAGAAACATATAAGTACTTTTCAC	167	
DB	592	TGTGGTGAACACCGTGGGCCGTGTACCATAATCGGCAGGAGACTGACCGGTACTTCTCTC	651	
QY	168	TTCCATTCTGTGTGGGCTCAAAAAAAGTATCAGTCAATTACATGAAACTCTGGGAGAG	227	
DB	652	TCCCTTTTGCAGTGGCCAAAAGTCTCTCGATATCCCACTACCAAGACGCTGAGCGAGG	711	
QY	228	CATTCAAGGGGTGAATTGGAAATTTAGTGGTCTGGATATTAATAATTAAAGATGATGTA	287	
DB	712	CGCTGAAGGAGTTCGAGCTGGAGTTCACTGGCTACGAGATGGAGTTCAAGAGCGACGCC	771	
QY	288	TGCCAGCCACTTACTGTGAAATTTGATTTAGATAAGAAAGAGAGATGTCATTTGTATATG	347	
DB	772	CCAAATCGGTCTCTGCATGGTCACTTTCGAGGGAGAGCGCCAAAGCATTCACCTATG	831	
QY	348	CCATAAAAAATCAATTACTGGTACCAAGTGPACATAGATGATTTACCAATATGGGGTATG	407	
DB	832	CCGTGAAGAACGAGTACTGGTACCAAAATGTACATCGATGGACTGCCCATTTTGGGGAAAG	891	
QY	408	TTTGGTCGGCTGATGAAATATGGAGAGATTAATCTTTTGGACCTATAAAAACTTGAAA	467	
DB	892	TGGTGAAGCGCAGGAGCGCGGATGGCAAGTACTATATCTTCACGCCAAGAAAGTTGACA	951	
QY	468	TAGGTTTTAATGAAATTCGAATTTGTTGATGTTAATCTAACTA	509	
DB	952	TCCGCTACATGATGCCAGCAAAATCGTGGATATCACCTTGACCA	993	

RESULT 14  
US-09-248-796A-6208  
; Sequence 6208, Application US/09248796A  
; Patent No. 6747137  
; GENERAL INFORMATION:  
; APPLICANT: Keith weinstock et al  
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICANS  
; TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS  
; FILE REFERENCE: 107196.132  
; CURRENT APPLICATION NUMBER: US/09/248,796A  
; CURRENT FILING DATE: 1999-02-12  
; PRIOR APPLICATION NUMBER: US 60/074,725  
; PRIOR FILING DATE: 1998-02-13  
; PRIOR APPLICATION NUMBER: US 60/096,409  
; PRIOR FILING DATE: 1998-08-13  
; NUMBER OF SEQ ID NOS: 28208  
; SEQ ID NO 6208  
; LENGTH: 726  
; TYPE: DNA  
; ORGANISM: Candida albicans  
US-09-248-796A-6208  
Query Match 7.3%; Score 132.8; DB 3; Length 726;

RESULT 13  
US-09-270-767-14715  
; Sequence 14715, Application  
; Patent No. 6703491  
; GENERAL INFORMATION:

Best Local Similarity 51.0%; Pred. No. 8e-25;  
Matches 367; Conservative 0; Mismatches 347; Indels 6; Gaps 2;  
QY 1031 GGAGGAAGAGATGGATTAAGCAGATGTTTATTTGGGGCAATCTCTATCCAGCTATGGTG 1090  
Db 13 GGTGGTGACAAATTTGGAAATTTGAATATGTTTTTGACACCAGTTTATAGTACCGAGGATTTTG 72  
QY 1091 TGTGGCAGTCCCTTCTTCATCAATTTTCATAGCAATTTATACCATGCTTTCAGAGCCATT 1150  
Db 73 TCTCTGGTTTTGTTGGTGTGTGAATTTCTTTAAATTTTCAGTACAATCTTCTGGTGTATT 132  
QY 1151 CTTTTTGGAAACAATGGTGGCCGTTTGTGCACTCTGTTTTTTTGTATTCTTCTCTCTAAAT 1210  
Db 133 CATATGGGGACAAATGTTGGCAATGCTTAATTTGGTCAATATATCGATTCATTAAGT 192  
QY 1211 CTTGTTGGTACAAATCTTGGCCGAATCTGTGAGGTGAGCCCAATCTTCTTGTGCGTGTG 1270  
Db 193 GTTATTTGGATCAATTTTAGCTAGTATAGCAATTAATATC---GGTACCAGTGAGAACT 249  
QY 1271 AATGCTGTGCTCGCTCTATACCGGAGAAATATGTTTCATGGAGCCTCGGTTATTGTT 1330  
Db 250 AATCAAAATTCAGACAAATCTCTACTCAACCATGATTTAAGTACTATCCCGGTAATG 309  
QY 1331 TGCCTGGTGGAAATTTTACCTTTTGGTTCATCTTTTATTTGAATGTTATTTCACTTTCACG 1390  
Db 310 TTTATTTCCGGAAATTTTCCCAATTTGGATCAATGCTGTGGAATGTTATTTATTTATCA 369  
QY 1391 TCTTTCTGGGCATATAAGATCTATTATGCTATGGCTTCATGATGCTGGTGGTGTATTC 1450  
Db 370 TCAATTTGGTTTTAATAAGATTTTATATGTTTGGATTTTATTTTCTGTTTCATATTA 429  
QY 1451 CTGTGCATTTGCTGCTGTGCTGACTATTGTGTCACATATTTTCTACTAAATCGAGAA 1510  
Db 430 ATGATTTTAACTAGTAGTTTAAATTAATTTTAAATGATTTTATTAATTTTATGTTTCAGAA 489  
QY 1511 GATTACCGGTGCAATGGCAAGTTTCTCTCTGCTGCTCAATCGCAATCTATGTTTAC 1570  
Db 490 AATTATAAATGCAATGGAATCAATATTGTTGGAGGAGTTGTGCAATTTATGTTATTT 549  
QY 1571 ATGATTTCTTTTACTACTATTATTTTCAAAACAAGATGATGGCTTATTTCAACATCA 1630  
Db 550 ATTCAATCAATTTTGTGCTGGTGTGA---AAAAATTTGGTGGATTTAGTTTCAATGTT 606  
QY 1631 TTTTACTTTGGATATATGGCGGTATTATAGCAGAGCTTTGGGATAATGTTGGAGCGATT 1690  
Db 607 TTATACAGTGGTATTACGCTGTGATTTCAATTTATGTTTCTTGTGTTGGATCAAT 666  
QY 1691 GGTACATGGGAACAAGTGCCTTTGTCGAAAAATCTATACATAATGTCAAAAATGACTAG 1750  
Db 667 GGATTTATAGTAGTTTAAATATTTGTCAGATTAATTTATGGTCAAAATTTAAAAATGATTAG 726

US-09-313-294A-2292  
Query Match 5.5%; Score 101; DB 3; Length 262;  
Best Local Similarity 68.4%; Pred. No. 1.2e-16;  
Matches 154; Conservative 0; Mismatches 70; Indels 1; Gaps 1;  
QY 1467 TCTGTGTGACTATTGTGTGCACATATTTTCTACTAAATGCAAGAAGATTACCGGTGGCAAT 1526  
Db 39 TCTCGGTCACTATTGTGGGTACTTTATTTCTTGTGAACGCCGAGAACTACCAATTGGCAAT 98  
QY 1527 GGACAAGTTTTTCTCTCTGCTGCATCAACTGCAATCTATGTTTACATGTTATTTCTTTACT 1586  
Db 99 GGAGTCTGTTTTTCTTCTGCAAGGTCAACCGCTCTGTAGGTGATCTGTACTCTCACTACT 158  
QY 1587 ACTATTTTTTCAAAACAAGATGATGGCTTATTTTCAACATCATTTTACTTTTGGATATA 1646  
Db 159 ACTACCATGTGAAGACAAAGATGTCCAGGCTTCTTCCAGACAAAGTTTCTTATTTGGCTACA 218  
QY 1647 TGGCGGTATTATTAGCACAGCCTTTGGGGATAATGTTGTGGAGCGATTG 1691  
Db 219 CGCTGATGTTCTGC-CTGGCCTAGGCATCTTTGTGGAGCTATTG 262

Search completed: March 10, 2006, 21:55:59  
Job time : 316.887 secs

RESULT 15  
US-09-313-294A-2292  
; Sequence 2292, Application US/09313294A  
; Patent No. 6476212  
; GENERAL INFORMATION:  
; APPLICANT: Lalgudi, Raghunath V.  
; APPLICANT: Ito, Laura Y.  
; APPLICANT: Sherman, Bradley K.  
; TITLE OF INVENTION: POLYNUCLEOTIDES AND POLYPEPTIDES DERIVED FROM CORN EAR.  
; FILE REFERENCE: PL-0017 US  
; CURRENT APPLICATION NUMBER: US/09/313,294A  
; CURRENT FILING DATE: 1995-05-14  
; NUMBER OF SEQ ID NOS: 7600  
; SOFTWARE: PERL Program  
; SEQ ID NO 2292  
; LENGTH: 262  
; TYPE: DNA  
; ORGANISM: Zea mays  
; FEATURES:  
; NAME/KEY: misc feature  
; OTHER INFORMATION: Incyte ID No. 6476212 700552439H1

Result No.	Query			DB	ID	Description
	Score	Match	Length			
1	1827	100.0	1827	8	US-10-755-466-3	Sequence 3, Appli-
2	1816.4	99.4	4508	3	US-09-814-353-21837	Sequence 21837, A
3	1816.4	99.4	4024	5	US-10-198-846-10005	Sequence 10005, A
4	1814	98.8	2072	8	US-10-755-466-1	Sequence 1, Appli
5	1804.4	98.8	3370	3	US-09-374-046A-25	Sequence 25, Appl
6	1804.4	98.8	3370	7	US-10-616-263-25	Sequence 25, Appl
7	1763	96.5	3076	3	US-09-913-582-29	Sequence 29, Appl
8	1763	96.5	3076	6	US-10-277-802-29	Sequence 29, Appl
9	1763	96.5	3076	8	US-10-896-972-29	Sequence 29, Appl
10	1725.4	94.4	3389	6	US-10-205-219-122	Sequence 122, App
11	1725.4	94.4	3389	9	US-10-956-157-2297	Sequence 2297, Ap
12	1725.4	94.4	3389	9	US-10-287-436A-335	Sequence 335, App
13	1024.6	56.1	6197	6	US-10-062-674-1697	Sequence 1697, Ap
14	726.4	39.8	1070	6	US-10-264-237-1414	Sequence 1414, Ap
15	646	35.4	1863	10	US-11-097-143-22277	Sequence 22277, A
16	629.6	34.5	1867	3	US-09-913-582-13	Sequence 13, Appl
17	629.6	34.5	1867	6	US-10-277-802-13	Sequence 13, Appl
18	629.6	34.5	1867	8	US-10-896-972-13	Sequence 140808,
19	583.6	31.9	2461	8	US-10-425-115-140808	Sequence 3465, Ap
20	581	31.8	2355	7	US-10-739-930-4365	Sequence 39405, A
21	573.6	31.4	1899	7	US-10-437-963-39405	Sequence 14430, A
22	570.8	31.2	2406	7	US-10-437-963-14430	Sequence 140919,
23	567.8	31.1	2698	8	US-10-425-115-140919	



















Qy	1115	TTCTAGGCCATTATTAATCACTGCTTCAAGAGCCATTCTTTTGGACAATAGTGGCGCGTT	1174
Db	1093	TTCTAGGCCATTATTAATCACTGCTTCAAGAGCCATTCTTTTGGACAATAGTGGCGCGTT	1152
Qy	1175	TGTTGGCATCTGTTTTTTTGTATTTCTTCCTCTAAATCTTGTTGGTCAAACTATTGGCGGA	1234
Db	1153	TGTTGGCATCTGTTTTTTTGTATTTCTTCCTCTAAATCTTGTTGGTCAAACTATTGGCGGA	1212
Qy	1235	AATCTGTCAAGTCAAGCCAACTTTCCTTGTGTGCTGCTCAATGCTGTGGCTCTGCTCTATACCG	1294
Db	1213	AATCTGTCAAGTCAAGCCAACTTTCCTTGTGTGCTGCTCAATGCTGTGGCTCTGCTCTATACCG	1272
Qy	1295	GAGAAAAATGTTTCATCGAGCCCTGCGGTATTGTTTGGCCCTGGGTGCAATTTTACCTTTT	1354
Db	1273	GAGAAAAATGTTTCATCGAGCCCTGCGGTATTGTTTGGCCCTGGGTGCAATTTTACCTTTT	1332
Qy	1355	GGTTCAATCTTTATGTAAATGTATTTCACTCTTCACGCTTTTCTGGGCAATAAAGATCTAT	1414
Db	1333	GGTTCAATCTTTATGTAAATGTATTTCACTCTTCACGCTTTTCTGGGCAATAAAGATCTAT	1392
Qy	1415	TATGCTCATGGCTTCATGATGCTGGTCTGGTTATCTGTGCATTTGTGCACTGTCTGTGTCG	1474
Db	1393	TATGCTCATGGCTTCATGATGCTGGTCTGGTTATCTGTGCATTTGTGCACTGTCTGTGTCG	1452
Qy	1475	ACTATTGTGTGCACATATTTTCTACTAAATGCAGAAGATTACCGGTGGCAATGGACAAGT	1534
Db	1453	ACTATTGTGTGCACATATTTTCTACTAAATGCAGAAGATTACCGGTGGCAATGGACAAGT	1512
Qy	1535	TTTCTCTCTCGTGCATCAACTGCAACTATGTGTTTCAATGATATCTCTTTTACTACTATTTT	1594
Db	1513	TTTCTCTCTCGTGCATCAACTGCAACTATGTGTTTCAATGATATCTCTTTTACTACTATTTT	1572
Qy	1595	TTCAAAACAAGATGATATGCTTATTTTCAAAACATCATTTTACTTTTGATATATGGCGGTA	1654
Db	1573	TTCAAAACAAGATGATATGCTTATTTTCAAAACATCATTTTACTTTTGATATATGGCGGTA	1632
Qy	1655	TTTAGCACAGCCTTGGGGATAAATGTGTGGAGCGGATGCGTTACATGGGAAACAAGTGCCCTT	1714
Db	1633	TTTAGCACAGCCTTGGGGATAAATGTGTGGAGCGGATGCGTTACATGGGAAACAAGTGCCCTT	1692
Qy	1715	GTCCGAAAATCTATACTAATGTGTAATTTGACTTAGAGACCCAGAGAAAACCTCGAACTTT	1774
Db	1693	GTCCGAAAATCTATACTAATGTGTAATTTGACTTAGAGACCCAGAGAAAACCTCGAACTTT	1752
Qy	1775	GGATCAATTTCTTTTTCATAGGGGTGGAACCTTGCAAGCAAAAAACAACAAAC	1827
Db	1753	GGATCAATTTCTTTTTCATAGGGGTGGAACCTTGCAAGCAAAAAACAACAAAC	1805

## RESULT 9

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US-10-896-972-29
; Sequence 29, Application US/10896972
; Publication No. US20050032168A1
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
; TITLE OF INVENTION: 17 Human Secreted Proteins
; FILE REFERENCE: PS723P1
; CURRENT APPLICATION NUMBER: US/10/896,972
; CURRENT FILING DATE: 2004-07-23
; PRIOR APPLICATION NUMBER: US/09/915,582
; PRIOR FILING DATE: 2001-07-27
; PRIOR APPLICATION NUMBER: PCT/US01/01431
; PRIOR FILING DATE: 2001-01-17
; PRIOR APPLICATION NUMBER: 60/179,065
; PRIOR FILING DATE: 2000-01-31
; PRIOR APPLICATION NUMBER: 60/180,628
; PRIOR FILING DATE: 2000-02-04
; PRIOR APPLICATION NUMBER: 60/231,968
; PRIOR FILING DATE: 2000-09-12
; NUMBER OF SEQ ID NOS: 97
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 29
; LENGTH: 3076

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585 TGAAATTTGAAGATCGATTTGACAAATATCTTGATCCGTC- TTTTTCACATCGGATT 643  
Db TGAATTTGAAGATCGATTTGACAAATATCTTGATCCGTCGCTTTTTCACATCGGATT 660  
644 CATTGGTTTTCAA-----TTTTCACATCTCTTCATGAT--GGTGATCTTCTTGGT--GGGCTT 696  
Db CATTGGTTTTCACATGTTTGTCACTCGTTCATGATGGTGATCTTCTTGGTGGGCTT 720  
697 AGTTTCAATGATTTTAATGAACA-----TTAAGAAAAAGATPATGCTCGGTACAGTAAAGA 753  
Db AGTTTCAATGATTTTAATGAACAATTAAGAAAAAGATPATGCTCGGTACAGTAAAGA 780  
754 GGAAGAAATGATGATATGATAGACACCTAGGAGATGAATATG--GATGAAAAAGGT 810  
Db GGAAGAAATGATGATATGATAGACACCTAGGAGATGAATATGATGATGGAAGAAAGGT 840  
811 GCAT--GGAGATGATTTAGACCAT--CAAGTCACCCACTGATATTTTCTCTCTGATGG 867  
Db GCATTTGAAGATGATTTAGACCATCAAGTCACCCACTGATATTTTCTCTCTGATGG 900  
868 TTCTGGATGTCAGATATTTGCTGTCTCTCATC--GTTATTTATTTGTCATGATAGAAG 926  
Db TTCTGGATGTCAGATATTTGCTGTCTCTCATCGTAAATGTTGTCATGATAGAAG 960  
927 ATTATATATAGAGAGGGATCAATGCTCAGTACAGCCATATTTGTCATGTC--TGCTAG 985  
Db ATTATATATAGAGAGGGATCAATGCTCAGTACAGCCATATTTGTCATGTCCTAGCTTACG 1020  
986 TCT--CCAGTGAATGTTATTTGAGGAGTCTGATCTAGACCAAGAGGAGGAGATG 1044  
Db TCTCCAGTGAATGTTATTTGAGGAGTCTGATCTAGACCAAGAGGAGGAGATG 1080  
1021 TCTCCAGTGAATGTTATTTGAGGAGTCTGATCTAGACCAAGAGGAGGAGATG 1080  
1045 GAT-----AAGACAGATGTTTATTTGGGGCAATTC-----TTATCCAGCTATGGTG 1090  
Db GCTATTAAGCCAGATGTTTATTTGGGGCAATTCCTTTTATTTCCCAAGCTAATGG 1140  
1091 TGTGGCACTGGCTTC-----TTATCAATTTTCATAGCCATTTATACATGCTT 1139  
Db GTTGTGTTGGCCCAATTCCTTTTTCATCAATTTTCATAGCCATTTATACATGCTT 1200  
1140 CAAGAGCCATTCCTTTTGGAAAC--AATGTCGCGCTTTGTCATCTG--TTTTTTTGTAT 1197  
Db CAAGAGCCATTCCTTTTGGAAACAAATGTCGCGCTTTGTCATCTGCTTTTGTAT 1260  
1198 TCTTCTCTAAATCTGTTGTCACATPACTTGGCCGAATCTGTAGGTCAGCCCAACTT 1257  
Db TCTTCTCTAAATCTGTTGTCACATPACTTGGCCGAATCTGTAGGTCAGCCCAACTT 1320  
1258 TCCTTGTGTCATGCTGTCCTC--GTGCTATACCGGAGA-----AAAAATGGTTTCATG 1312  
Db TCCTTGTGTCATGCTGTCCTCCTGCTATACCGGAGAACACACAGATGGTACATG 1380  
1313 GAGCTCGGTTATTTGCTGGGGGAATTTTACCTTTGCTCAATCTTTATTGAA 1372  
Db GAGCTCGGTTATTTGCTGGGGGAATTTTACCTTTGCTCAATCTTTATTGAA 1440  
1373 ATGTTATTTCACTTTTCAGCTCTTTCTGGGCATATAAGATCTATTATGCTATGGCTTCATG 1432  
Db ATGTTATTTCACTTTTCAGCTCTTTCTGGGCATATAAGATCTATTATGCTATGGCTTCATG 1500  
1433 ATGCTGTGCTGGTTATCTGTGTCATTTGTCATGCTGTCGTCATTTGTCGACATAT 1492  
Db ATGCTGTGCTGGTTATCTGTGTCATTTGTCATGCTGTCGTCATTTGTCGACATAT 1560  
1493 TTCTCTAATGACAGAGATTAACCGGTGGCAATGGACAAGTTTCTCTCTGCTGTCATCA 1552  
Db TTCTCTAATGACAGAGATTAACCGGTGGCAATGGACAAGTTTCTCTCTGCTGTCATCA 1620  
1553 ACTGCAATCTATGTTTACATGATTTCTTTTACTACTATTTTTT-----CAAAACAAA 1605  
Db ACTGCAATCTATGTTTACATGATTTCTTTTACTACTATTTTTTCTGGAACAAAGATG 1680

1606 GATGATGCTTATTTCAACATCATTTTACTTTGGATATATGGC--GGTATTTAGCACA 1663  
Db TATGTCCTTATTTGTCAACATCTTATTTTACATTTGGATATATGGCTGTATATAGCACA 1740  
1664 GCCTTGGGATATATGTTG---GAGCGATTTGTTACATGGGAACAAGTGCCTTTTGTCCG 1719  
Db GTCTTGGGATATATGTTGTTGGAGCGATATGTTTACATGGGAACAAGTGCCTTTTGTCCG 1800  
1720 AAAATCTATCTAATGTTGAATAATGACTAGACCCAGAACAACTTGGAACTTT--GGAT 1778  
Db AAAATCTATCTAATGTTGAATAATGACTAGACCCAGAACAACTTGGAACTTTTGGAT 1860  
1779 CAATTTCTTTTTCATAGGGGT--GGAATTTGCACAGCAAAAAACAAC 1827  
Db CAATTTCTTTTTCATAGGGGTGGGAATTTGCACAGCAAAAAACAAC 1910

RESULT 14  
US-10-264-237-1414  
; Sequence 1414, Application US/10264237  
; Publication No. US20040009491A1  
; GENERAL INFORMATION:  
; APPLICANT: Birse et al.  
; TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies  
; FILE REFERENCE: P4131P1  
; CURRENT APPLICATION NUMBER: US/10/264,237  
; CURRENT FILING DATE: 2002-10-04  
; PRIOR APPLICATION NUMBER: PCT/US01/16450  
; PRIOR FILING DATE: 2001-05-18  
; PRIOR APPLICATION NUMBER: US 60/205,515  
; PRIOR FILING DATE: 2000-05-19  
; NUMBER OF SEQ ID NOS: 2876  
; SOFTWARE: Patent In Ver. 3.1  
; SEQ ID NO 1414  
; LENGTH: 1070  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; FEATURE:  
; NAME/KEY: misc\_feature  
; LOCATION: (34)..(34)  
; OTHER INFORMATION: n equals a,t,g, or c  
; FEATURE:  
; NAME/KEY: misc\_feature  
; LOCATION: (40)..(40)  
; OTHER INFORMATION: n equals a,t,g, or c  
; FEATURE:  
; NAME/KEY: misc\_feature  
; LOCATION: (525)..(525)  
; OTHER INFORMATION: n equals a,t,g, or c  
; FEATURE:  
; NAME/KEY: misc\_feature  
; LOCATION: (529)..(529)  
; OTHER INFORMATION: n equals a,t,g, or c  
; FEATURE:  
; NAME/KEY: misc\_feature  
; LOCATION: (557)..(557)  
; OTHER INFORMATION: n equals a,t,g, or c  
; FEATURE:  
; NAME/KEY: misc\_feature  
; LOCATION: (837)..(837)  
; OTHER INFORMATION: n equals a,t,g, or c  
; FEATURE:  
; NAME/KEY: misc\_feature  
; LOCATION: (912)..(912)  
; OTHER INFORMATION: n equals a,t,g, or c  
; FEATURE:  
; NAME/KEY: misc\_feature  
; LOCATION: (956)..(956)  
; OTHER INFORMATION: n equals a,t,g, or c  
; FEATURE:  
; NAME/KEY: misc\_feature  
; LOCATION: (965)..(966)  
; OTHER INFORMATION: n equals a,t,g, or c







121 ACATCTCCAGTGAATGGGTAATTTTGGG

QY 1043 TGGATAAAGCAGATGTTTATTTGGGGCATTCCTTATCCAGCTATGGTGTGGCACTGCC 1102  
DB 181 TGGATAAAGCAGATGTTTATTTGGGGCATTCCTTATCCAGCTATGGTGTGGCACTGCC 240  
QY 1103 TTCTTCATCAATTTATAGCCATTTATACCATGCTTCAAGAGCCATTCCTTTTGGAAACA 1162  
DB 241 TTCTTCATCAATTTATAGCCATTTATACCATGCTTCAAGAGCCATTCCTTTTGGAAACA 300  
QY 1163 ATGGTGGCCGTTGTTGGCATCTGTTTTTTTGTATATCTTCTTAAATCTTGTGGTACA 1222  
DB 301 ATGGTGGCCGTTGTTGGCATCTGTTTTTTTGTATATCTTCTTAAATCTTGTGGTACA 360  
QY 1223 ATACTTGGCCGAATCTGTAGGTCAGGCCAACTTCTTGTGCTGCAATGCTGTGCT 1282  
DB 361 ATACTTGGCCGAATCTGTAGGTCAGGCCAACTTCTTGTGCTGCAATGCTGTGCT 420  
QY 1283 CGTCTATACCGAGAAAATGTTTCATGGAGCTCGGGTTATTTGTTGCTGGGTGGA 1342  
DB 421 GGTCTATACCGAGAAAATGTTTCATGGAGCTCGGGTTATTTGTTGCTGGGTGGA 480  
QY 1343 ATTTTACCTTTTGGTTCAATCTTTTATGAAATGTTTATTCATCTTCAAGCTCTTCTGGGCA 1402  
DB 481 ATTTTACCTTTTGGTTCAATCTTTTATGAAATGTTTATTCATCTTCAAGCTCTTCTGGGCA 540  
QY 1403 TATAAGATCTA-TTATGCTATGGCTTCAT 1431  
DB 541 TACAAGACCCACTTATGCTATGGCTTTAT 570

## RESULT 2

US-11-128-061-4967  
; Sequence 4967, Application US/11128061  
; Publication No. US2006003958A1  
; GENERAL INFORMATION:  
; APPLICANT: Melville, Mark W.  
; APPLICANT: Charlebois, Timothy S.  
; APPLICANT: Mounts, William M.  
; APPLICANT: Hann, Louane E.  
; APPLICANT: Sinacore, Martin S.  
; APPLICANT: Leonard, Mark W.  
; APPLICANT: Brown, Eugene L.  
; APPLICANT: Miller, Christopher P.  
; TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES RELATED TO OLIGONUCLEOTIDE ARRAYS  
; FILE REFERENCE: 01997.027701  
; CURRENT APPLICATION NUMBER: US/11/128,061  
; PRIOR FILING DATE: 2005-05-11  
; PRIOR APPLICATION NUMBER: US 60/570,425  
; PRIOR FILING DATE: 2004-05-11  
; NUMBER OF SEQ ID NOS: 7285  
; SOFTWARE: PatentIn version 3.3  
; SEQ ID NO 4967  
; LENGTH: 570  
; TYPE: DNA  
; ORGANISM: Cricetulus griseus  
US-11-128-061-4967

Query Match 26.4%; Score 482.8; DB 12; Length 570;  
Best Local Similarity 91.6%; Pred. No. 4.8e-102;  
Matches 522; Conservative 0; Mismatches 47; Indels 1; Gaps 1;  
QY 863 ATTGGTTCTGGATGTCAGATATTTGCTGTCTCTCATCGTTATTTGTTGCAATGATA 922  
DB 1 ATTGGTTCTGGATGTCAGATATTTGCTGTCTCTCATCGTTATTTGTTGCAATGATA 60  
QY 923 GAAGATTTATATCTAGAGGGGATCAATGCTCAGTACAGCCATATTTGCTATGCTGCT 982  
DB 61 GAGGATTTATATACAGAGAGGGGATCAATGCTCAGTACAGCCATATTTGCTATGCTGCT 120  
QY 983 ACGTCTCCAGTCAATGTTTATTTTGGAGGAAGTCTGTATGCTAGACAAGGAGGAGAGA 1042  
DB 121 ACATCTCCAGTCAATGTTTATTTTGGAGGAAGTCTGTATGCTAGACAAGGAGGAGAGA 180

QY 1043 TGGATAAAGCAGATGTTTATTTGGGGCATTCCTTATCCAGCTATGGTGTGGCACTGCC 1102  
DB 181 TGGATAAAGCAGATGTTTATTTGGGGCATTCCTTATCCAGCTATGGTGTGGCACTGCC 240  
QY 1103 TTCTTCATCAATTTATAGCCATTTATACCATGCTTCAAGAGCCATTCCTTTTGGAAACA 1162  
DB 241 TTCTTCATCAATTTATAGCCATTTATACCATGCTTCAAGAGCCATTCCTTTTGGAAACA 300  
QY 1163 ATGGTGGCCGTTGTTGGCATCTGTTTTTTTGTATATCTTCTTAAATCTTGTGGTACA 1222  
DB 301 ATGGTGGCCGTTGTTGGCATCTGTTTTTTTGTATATCTTCTTAAATCTTGTGGTACA 360  
QY 1223 ATACTTGGCCGAATCTGTAGGTCAGGCCAACTTCTTGTGCTGCAATGCTGTGCT 1282  
DB 361 ATACTTGGCCGAATCTGTAGGTCAGGCCAACTTCTTGTGCTGCAATGCTGTGCT 420  
QY 1283 CGTCTATACCGAGAAAATGTTTCATGGAGCTCGGGTTATTTGTTGCTGGGTGGA 1342  
DB 421 GGTCTATACCGAGAAAATGTTTCATGGAGCTCGGGTTATTTGTTGCTGGGTGGA 480  
QY 1343 ATTTTACCTTTTGGTTCAATCTTTTATGAAATGTTTATTCATCTTCAAGCTCTTCTGGGCA 1402  
DB 481 ATTTTACCTTTTGGTTCAATCTTTTATGAAATGTTTATTCATCTTCAAGCTCTTCTGGGCA 540  
QY 1403 TATAAGATCTA-TTATGCTATGGCTTCAT 1431  
DB 541 TACAAGACCCACTTATGCTATGGCTTTAT 570

## RESULT 3

US-11-128-049-1325  
; Sequence 1325, Application US/11128049  
; Publication No. US20060010513A1  
; GENERAL INFORMATION:  
; APPLICANT: Melville, Mark W.  
; APPLICANT: Charlebois, Timothy S.  
; APPLICANT: Mounts, William M.  
; APPLICANT: Hann, Louane E.  
; APPLICANT: Sinacore, Martin S.  
; APPLICANT: Leonard, Mark W.  
; APPLICANT: Brown, Eugene L.  
; APPLICANT: Miller, Christopher P.  
; TITLE OF INVENTION: OLIGONUCLEOTIDE ARRAYS TO MONITOR GENE EXPRESSION AND METHODS FOR  
; FILE REFERENCE: 01997.027700  
; CURRENT APPLICATION NUMBER: US/11/128,049  
; PRIOR FILING DATE: 2005-05-11  
; PRIOR APPLICATION NUMBER: US 60/570,425  
; PRIOR FILING DATE: 2004-05-11  
; NUMBER OF SEQ ID NOS: 7285  
; SOFTWARE: PatentIn version 3.3  
; SEQ ID NO 1325  
; LENGTH: 570  
; TYPE: DNA  
; ORGANISM: Cricetulus griseus  
US-11-128-049-1325

Query Match 26.4%; Score 482.8; DB 12; Length 570;  
Best Local Similarity 91.6%; Pred. No. 4.8e-102;  
Matches 522; Conservative 0; Mismatches 47; Indels 1; Gaps 1;  
QY 863 ATTGGTTCTGGATGTCAGATATTTGCTGTCTCTCATCGTTATTTGTTGCAATGATA 922  
DB 1 ATTGGTTCTGGATGTCAGATATTTGCTGTCTCTCATCGTTATTTGTTGCAATGATA 60  
QY 923 GAAGATTTATATCTAGAGGGGATCAATGCTCAGTACAGCCATATTTGCTATGCTGCT 982  
DB 61 GAGGATTTATATACAGAGAGGGGATCAATGCTCAGTACAGCCATATTTGCTATGCTGCT 120  
QY 983 ACGTCTCCAGTCAATGTTTATTTTGGAGGAAGTCTGTATGCTAGACAAGGAGGAGAGA 1042  
DB 121 ACATCTCCAGTCAATGTTTATTTTGGAGGAAGTCTGTATGCTAGACAAGGAGGAGAGA 180

QY 1043 TGGATAAGCAGATGTTTATTTGGGCAATTCCTTATCCAGCTATGGTGTGGCACTGCC 1102  
DB 181 TGGATAAGCAGATGTTTATTTGGGCAATTCCTTATCCAGCTATGGTGTGGCACTGCC 240  
QY 1103 TTCTTCATCAATTCATAGCATTATTTACATGCTTCAAGAGCCATTCCTTTTGGAAACA 1162  
DB 241 TTCTTCATCAATTCATAGCATTATTTACATGCTTCAAGAGCCATTCCTTTTGGAAACA 300  
QY 1163 ATGGTGGCCGTTGTTGTCATCTGTTTTTGTATTTCTTCTCTAAATCTTGTGTGTA 1222  
DB 301 ATGGTGGCCGTTGTTGTCATCTGTTTTTGTATTTCTTCTCTAAATCTTGTGTGTA 360  
QY 1223 ATACTTGGCCGAATCTGTGAGTCAGCCCAACTTCTTGTGTGTCATGCTGTGCCCT 1282  
DB 361 ATACTTGGCCGAATCTGTGAGTCAGCCCAACTTCTTGTGTGTCATGCTGTGCCCT 420  
QY 1283 CGTCTATACCGGAGAAAAATGTTTCATGGAGCCTGCGGTATTTGTCCTGGGTGGA 1342  
DB 421 GGTCTATCCAGACAAATATGTTTATGATCTCTGATTTATCGTTCGCTGTAGCA 480  
QY 1343 ATTTTACCTTTTGGTTCATCTTTATTTGAATGTTATTCATCTTCAAGTCTTCTGGGCA 1402  
DB 481 ATTTTACCTTTTGGTTCATCTTTATTTGAATGTTATTCATCTTCAAGTCTTCTGGGCA 540  
QY 1403 TATAAGATCTA-TTATGTCATGCTTCAT 1431  
DB 541 TACAAGACCCACTTATGTCATGGCTTTAT 570

## RESULT 4

US-11-128-049-4967  
; Sequence 4967, Application US/11128049  
; Publication No. US20060010513A1  
; GENERAL INFORMATION:  
; APPLICANT: Melville, Mark W.  
; APPLICANT: Charlebois, Timothy S.  
; APPLICANT: Mounts, William M.  
; APPLICANT: Hann, Louane E.  
; APPLICANT: Sinacore, Martin S.  
; APPLICANT: Leonard, Mark W.  
; APPLICANT: Brown, Eugene L.  
; APPLICANT: Miller, Christopher P.  
; TITLE OF INVENTION: OLIGONUCLEOTIDE ARRAYS TO MONITOR GENE EXPRESSION AND METHODS FOR  
; FILE REFERENCE: 01997.027700  
; CURRENT APPLICATION NUMBER: US/11/128,049  
; CURRENT FILING DATE: 2005-05-11  
; PRIOR APPLICATION NUMBER: US 60/570,425  
; PRIOR FILING DATE: 2004-05-11  
; NUMBER OF SEQ ID NOS: 7285  
; SOFTWARE: Patent in version 3.3  
; SEQ ID NO 4967  
; LENGTH: 570  
; TYPE: DNA  
; ORGANISM: Cricetulus griseus  
US-11-128-049-4967

Query Match 26.4%; Score 482.8; DB 12; Length 570;  
Best Local Similarity 91.6%; Pred. No. 4.8e-102;  
Matches 522; Conservative 0; Mismatches 47; Indels 1; Gaps 1;

QY 863 ATTGGTCTGATGTGATGATTTGCTGCTCTCATGCTTATTTATTTGTCATGATA 922  
DB 1 ATTGGTCTGATGTGATGATTTGCTGCTCTCATGCTTATTTATTTGTCATGATA 60  
QY 923 GAGATTTATATCTAGAGGGATCAATCTCAGTACAGCCATTTTGTCTATGCTGCT 982  
DB 61 GAGATTTATATACAGAGAGGGATCAATGCTCAGTACAGCCATTTTGTCTATGCTGCT 120  
QY 983 ACCTCTCCAGTGAATGTTATTTTGGAGGAAGTCTGTATCTAGACAAGAGGAGAGA 1042  
DB 121 ACATCTCCAGTGAATGTTATTTTGGAGGAAGTCTGTATCTAGACAAGAGGAGAGA 180

QY 1043 TGGATAAGCAGATGTTTATTTGGGCAATTCCTTATCCAGCTATGGTGTGGCACTGCC 1102  
DB 181 TGGATAAGCAGATGTTTATTTGGGCAATTCCTTATCCAGCTATGGTGTGGCACTGCC 240  
QY 1103 TTCTTCATCAATTCATAGCATTATTTACATGCTTCAAGAGCCATTCCTTTTGGAAACA 1162  
DB 241 TTCTTCATCAATTCATAGCATTATTTACATGCTTCAAGAGCCATTCCTTTTGGAAACA 300  
QY 1163 ATGGTGGCCGTTGTTGTCATCTGTTTTTGTATTTCTTCTCTAAATCTTGTGTGTA 1222  
DB 301 ATGGTGGCCGTTGTTGTCATCTGTTTTTGTATTTCTTCTCTAAATCTTGTGTGTA 360  
QY 1223 ATACTTGGCCGAATCTGTGAGTCAGCCCAACTTCTTGTGTGTCATGCTGTGCCCT 1282  
DB 361 ATACTTGGCCGAATCTGTGAGTCAGCCCAACTTCTTGTGTGTCATGCTGTGCCCT 420  
QY 1283 CGTCTATACCGGAGAAAAATGTTTCATGGAGCCTGCGGTATTTGTCCTGGGTGGA 1342  
DB 421 GGTCTATCCAGACAAATATGTTTATGATCTCTGATTTATCGTTCGCTGTAGCA 480  
QY 1343 ATTTTACCTTTTGGTTCATCTTTATTTGAATGTTATTCATCTTCAAGTCTTCTGGGCA 1402  
DB 481 ATTTTACCTTTTGGTTCATCTTTATTTGAATGTTATTCATCTTCAAGTCTTCTGGGCA 540  
QY 1403 TATAAGATCTA-TTATGTCATGCTTCAT 1431  
DB 541 TACAAGACCCACTTATGTCATGGCTTTAT 570

## RESULT 5

US-11-000-688-1309  
; Sequence 1309, Application US/11000688  
; Publication No. US20050287544A1  
; GENERAL INFORMATION:  
; APPLICANT: BERTUCCI, Francois  
; APPLICANT: HOULGAITE, Remi  
; APPLICANT: BIRNEAUM, Daniel  
; TITLE OF INVENTION: GENE EXPRESSION PROFILING OF COLON CANCER WITH DNA ARRAYS  
; FILE REFERENCE: 1423-R-03  
; CURRENT APPLICATION NUMBER: US/11/000,688  
; CURRENT FILING DATE: 2004-12-01  
; PRIOR APPLICATION NUMBER: US 60/525,987  
; PRIOR FILING DATE: 2003-12-01  
; NUMBER OF SEQ ID NOS: 1596  
; SOFTWARE: Patent in version 3.2  
; SEQ ID NO 1309  
; LENGTH: 2391  
; TYPE: DNA  
; ORGANISM: Artificial Sequence  
; FEATURE:  
; OTHER INFORMATION: Description of Artificial sequences: primer  
; FEATURE:  
; NAME/KEY: misc feature  
; LOCATION: (1)-(2391)  
; OTHER INFORMATION: transmembrane 9 superfamily member 2 (TMS9F2)  
; OTHER INFORMATION: gene.  
US-11-000-688-1309

Query Match 12.5%; Score 227.6; DB 12; Length 2391;  
Best Local Similarity 51.9%; Pred. No. 1.2e-42;  
Matches 596; Conservative 0; Mismatches 534; Indels 18; Gaps 3;

QY 609 AATATCTTGATCCGCTCTTTTCAACATCGGATTCATTTGGTTTCAATTTTCACTCCT 668  
DB 996 ACTATATCTGAGTCTATGCTCTATACCCACATTCAGTGTGTAGCATTTAGCAATCC 1055  
QY 669 TCATGATGATGATCTCTTGTGGCTTAGTTTCAATGATTTTATGAGAACATTAGAA 728  
DB 1056 TGGTCATTTCTCTCTTATCTGGAATGGTAGTAATGATTTAGTACGACACTGCACA 1115  
QY 729 AAGATTTATGCTCGGTACAGTAAAGAGGAAGAAATGGATGATATGATAGAGACTAGGAG 788

Db 1116 AAGATATTGCTAGATATAATCAGATGGACTCTACGGAGATGCCAG-----G 1163  
Qy 789 ATGAATATGGATGGAACAGGTGATGAGATGATTTAGAACATCAAGTCACCACTGA 848  
Db 1164 AAGAATTTGGCTGGAACCTTTCATGTGTATATATTCGCTCTCCAGAAAGGGATGC 1223  
Qy 849 TATTTTCTCTCTGATGTTCTCGATGTCAGATATTTGCTGTCTCTCATCGTTATTA 908  
Db 1224 TGCTATCAGTCTTTCTAGATCCGGGACACAGATTTTAAATATGACCTTTGTGACTCTAT 1283  
Qy 909 TTGTGCAATGATAGAGATTTATATCTAGAGGGGATCAATCTCAGTAC---AGCCA 965  
Db 1284 TTTTCGCTTGCTGGATTTTGTTCACCTGCCACCGAGGAGCGCTGATGAGTGTGCTG 1343  
Qy 966 TATTTGCTATGCTGTACGTCCTCAGTGAATGTTATTTTGGAGGAAGTCTGTATGCTA 1025  
Db 1344 TGGTCTGTGGGTGCTGCTGGGACCCCTGCGAGCTATGTTGCTGCCAGATTTCTATAAGT 1403  
Qy 1026 GACAAGGAGGAAGAGATGGATGAAGCAGATGTTTATTTGGGGCAATCCTTATCCAGCTA 1085  
Db 1404 CTTTGGAGGTGAGAGTGGGAAACAAATGTTTATTAACATCAITTTCTTTGTCTGGGA 1463  
Qy 1086 TGGTGTGTGGCAGTCTCTTCAATCAATTTTCATAGCCATTTATTAACCATGCTTCAAGAG 1145  
Db 1464 TTGTATTTGCTGACTTCTTATATATGATCTGATCTCTGCGGAGAGGATCTTCAGCAG 1523  
Qy 1146 CCATTCCTTTTGGAAACAAATGGTGGCCGTTTGTGCTATCTGTTTGTATTTCTTCTC 1205  
Db 1524 CTATTCCTTTTGGACACAGTGGTCCCATATGTTGCCCTTGTGTTCTGCATATCTGTGCTC 1583  
Qy 1206 TAAATCTTGTGGTACATACTTGGCGGAATCTGTCAGGTCAGCCCAACTTCTTCTGTC 1265  
Db 1584 TGAGTTTATTTGGTGATACCTTTGGTTTAAAGAAATGCCATGAACAC---CCAGTTC 1640  
Qy 1266 GTGTCAATGTGTGCTGCTCTATACCGGAGAAATGTTTCATGGAGCTGCGGTTA 1325  
Db 1641 GAACCAATCAGATTTCCACGTCAGATTCCTGAACAGTCTGTTCTACACGAAGCCCTTGCTG 1700  
Qy 1326 TTGTTTCCCTGGTGGAAATTTTACCTTTTGGTTCAATCTTTATTTGAATGATTTTCACT 1385  
Db 1701 GTATTATATCGGAGGAGATTTTGGCCCTTTGGCTGCATCTTTATACAACTTTTCTTCA 1760  
Qy 1386 TCAGTCTTCTTGGGCATATAAGATCTATTATGCTATGCTTTCATGATGCTGTGCTGG 1445  
Db 1761 TGAATAGTATTTGTTCACACAGATGATTTACATGTTTGGCTTCTTATTTCTGTGTTA 1820  
Qy 1446 TTATCTGTGATGTGACTGCTGTGTGACTATTTGTGTCACATATTTTCTACTAAATG 1505  
Db 1821 TCAATTTGGTTATTAACCTGTTCTGAAGCAACTATACCTTCTTTGCTATTTCCACCTATGT 1880  
Qy 1506 CAGAGATTAACCGTGGCAATGGACAAGTTTCTCTGCTGCTGATCACTGCAATCTATG 1565  
Db 1881 CAGAGGATTAATCAATGGCAATGGCGTTCAITTCCTTACGAGTGGCTTTTACTGCAATTA 1940  
Qy 1566 TTTTACATGATTCCTTTTACTACTATTTTCAAAACAAAGATGATGCTTATTTTCAAA 1625  
Db 1941 TCTTAATCTATGAGTACACTACTCTTTTCAAAATCTGCAGATCAGCGGAACAGCAGCA 2000  
Qy 1626 CATCAITTTTACTTTTGGATATATGGCGTATTTTAGCACAGCGCTTTGGGATATATGTTGGAG 1685  
Db 2001 CAATTCGTACTTTTGGTTATACCATGATATGTTGTTTGTATCTTCTTTTACAGGAA 2060  
Qy 1686 CGATGTTTACATGGGAACAAGTGCCTTTTGTCCGAAAATCTACTAATGTTGAAATG 1745  
Db 2061 CAATTTGCTCTTTTGTGATGCTTTTGTGTTTGTGTACCAAAATATACAGTGTGTGAGGTTG 2120  
Qy 1746 ACTAGAGA 1753  
Db 2121 ACTGAAGA 2128

RESULT 6

US-11-072-512-1699

; Sequence 1699, Application US/11072512  
; Publication No. US20060029945A1  
; GENERAL INFORMATION:  
; APPLICANT: ISOGAI, TAKAO  
; APPLICANT: SUGIYAMA, TOMOYASU  
; APPLICANT: OTSUKI, TETSUJI  
; APPLICANT: WAKAMATSU, AI  
; APPLICANT: SATO, HIROYUKI  
; APPLICANT: ISHII, SHIZUKO  
; APPLICANT: YAMAMOTO, JUN-ICHI  
; APPLICANT: ISONO, YUUKO  
; APPLICANT: HIO, YURI  
; APPLICANT: OTSUKA, KAORU  
; APPLICANT: NAGAI, KEIICHI  
; APPLICANT: IRIE, RYOTARO  
; APPLICANT: TAMECHIKA, ICHIRO  
; APPLICANT: SEKI, NAOHICO  
; APPLICANT: YOSHIKAWA, TSUTOMU  
; APPLICANT: OTSUKA, MOTOKYUKI  
; APPLICANT: NAGAHARI, KENJI  
; APPLICANT: MASUHO, YASUHIKO  
; TITLE OF INVENTION: Novel full length cDNA  
; FILE REFERENCE: 084335-0191  
; CURRENT APPLICATION NUMBER: US/11/072,512  
; CURRENT FILING DATE: 2005-03-07  
; PRIOR APPLICATION NUMBER: US 60/350,978  
; PRIOR FILING DATE: 2002-01-25  
; PRIOR APPLICATION NUMBER: JP 2001-379298  
; PRIOR FILING DATE: 2001-11-05  
; NUMBER OF SEQ ID NOS: 4096  
; SOFTWARE: PatentIn Ver. 2.1  
; SEQ ID NO 1699  
; LENGTH: 1878  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; US-11-072-512-1699  
Query Match 12.4%; Score 226; DB 9; Length 1878;  
Best Local Similarity 51.8%; Pred. No. 2,6e-42;  
Matches 595; Conservative 0; Mismatches 535; Indels 18; Gaps 3;  
Qy 609 AATATCTTGATCCGTCCTTTTCAACATCGGATTCATTTGGTTTCAATTTTCAATCTCT 668  
Db 693 ACTATATCTGGAGTCTATGCTCATPACCCACATTCAGTGGTTTAGCAATTATGAATTC 752  
Qy 669 TATGATGTGATCTTCTTGGTGGCTTAGTTTCAATGATTTTATGAGACATTAGAA 728  
Db 753 TGTCTATTTCTTCTTCTTATCTGGAATGTTAGTATGATTTTACGGACACTGCACA 812  
Qy 729 AAGATTATCTCGGTACAGTAAAGAGGAAGAAATGATGATATGATAGAGACCTAGGAG 788  
Db 813 AAGATATCTAGATATAATCAGATGGATCTACGGAGATGCCAG-----G 860  
Qy 789 ATGAATATCGATGGAAACAGGTGCTAGTGGAGATGATTTTAGACCATCAAGTCACCACTGA 848  
Db 861 AAGATTTGCTGGAACTTGTTCATGTTGATATATTCGCTCTCCAGAAAGGGATGC 920  
Qy 849 TATTTTCTCTCTGATGTTGTTCTGGATGTCAGATATTTGCTGTCTCTCATCGTTATTA 908  
Db 921 TGCTATCAGTCTTCTTAGGATCCGGGACACAGATTTTAAATATGACCTTTGTGACTCTAT 980  
Qy 909 TTGTGCAATGATAGAAGATTTATATCTAGAGGGGATCAATGCTCAGTAC---AGCCA 965  
Db 981 TTTTCCGCTTGGATTTTGTACCTGCCAACCGAGGAGCGCTGATGACGTTGCTG 1040  
Qy 966 TATTTGCTATGCTGCTACGTCCTCCAGTGAATGGTTTATTTTGGAGGAAGTCTGTATGCTA 1025  
Db 1041 TGCTCTGTGGTGTGCTGGGCACCCCTGCGAGCTATGTTGCTGCCAGATTTCTATAAGT 1100  
Qy 1026 GACAGGAGGAGGAGATGGATAAGCAGATGTTTATTTGGGCAITTCCTTATCCAGCTA 1085  
Db 1101 CCTTTGGAGGTGAGAGTGGAAACAAATGTTTATTAACATCAITTTCTTGTCTCGGGA 1160

1086 TGGTGTGGCACTGCTTCTTCATCAATTTTCATAGCCATTTATTTACCATGCTTCAAGAG 1145  
1161 TTGTATTTGCTGACTTCTTTTATAATGAATCTGATCTCTCGGGAGAGGATCTTTCAGCAG 1220  
1146 CCATTCCTTTTGGAAACAATGGTGGCCGTTTGTGGCATCTGTTTTTTTGTATTCTTCTTC 1205  
1221 CTATTCCTTTTGGACACATGGTGGCCATATGGCCCTTTGGTTCTGCATATCTGTGCTTC 1280  
1206 TAAATCTTGTGTGTACAACTTGTGGCGAAATCTGTGAGTCAGCCCAACTTTTCTTGTGTC 1265  
1281 TGACGTTTATTTGTGTGATCTTTTGGTTTAAAGAAATGCCATTTGAACAC---CCAGTTTC 1337  
1266 GTGTCATGCTGTGCTGCTGCTTATACCGGAGAAATAATGGTTTCATGGAGCCCTGCGGTTA 1325  
1338 GAACCAATCAGATTCCACGTCAGATTCTCTGAACAGTGTCTACAGAAAGCCCTTGCCTG 1397  
1326 TTGTTTGGCTGGGTGGAAATTTTACCTTTTGGTTTCAATCTTTTATTTGAAATGTAATTTTCATCT 1385  
1398 GTATTATCATGGAGGGAATTTTGGCCCTTTGGCTGTCATCTTTTATACAACTTTTCTTCAATTC 1457  
1386 TCACGCTTTTCTGGGCATATAAGATCTATTAATCTTATGCTTATGCTTCATGATGCTGGTCTGG 1445  
1458 TGAATAGTATTTGGTCAACACAGATGTAATACATGTTTGGCTTCTTATTTCTGGTGTAA 1517  
1446 TTATCTGTGCAATTTGTGCTGTGTGACTATTGTGTGCATATTTTCTACTAAATG 1505  
1518 TCATTTTGGTTATTTACTGTTCTGTAAGCAACTATCTTCTTGTCTATTTTCCACTATGTG 1577  
1506 CAGAAGATTACCGGTGGCAATGACAAAGTTTCTCTGCTGTCATCAACTGCAATCTATG 1565  
1578 CAGAGATTATCATTTGGCAATGGGTTTCACTCTTAGAGTGGCTTTTACTGCAATTTAT 1637  
1566 TTTACATGATTTCTTTTACTACTATTTTTCAAAACAAAGATGTAAGCTTATTTTCAAA 1625  
1638 TCTTAATCTATGTCAGTACACTACTTCTTTTCAAACTGACAGATCAACGGGAACAGACG 1697  
1626 CATCATTTTACTTTGATATATGGGTTATTTAGCACAGCTTGGGATAATGTGTGGAG 1685  
1698 CAATTCGTACTTTGGTTATACCATGATATGTTTGTATGTTTGTATGTTTCTTTTACAGGAA 1757  
1686 CGATTTGGTTACATGGGAACAAGTGGCTTTTGTCCGAAAAATCTATCTAATATGTGAAAAAT 1745  
1758 CAATTTGGCTTTTGTATGCTTTTGGTTTGTATACCAATATACATGTTGGTGAAGTTG 1817  
1746 ACTAGAGA 1753  
1818 ACTGAAGA 1825

## RESULT 7

US-11-240-769-50  
; Sequence 50, Application US/11240769  
; Publication No. US20060036089A1  
; GENERAL INFORMATION:  
; APPLICANT: Soppet et al.  
; TITLE OF INVENTION: 33 Human Secreted Proteins  
; FILE REFERENCE: P2037P1C2  
; CURRENT APPLICATION NUMBER: US/11/240,769  
; CURRENT FILING DATE: 2005-10-03  
; PRIOR APPLICATION NUMBER: 09/997,131  
; PRIOR FILING DATE: 2001-11-30  
; PRIOR APPLICATION NUMBER: 09/628,508  
; PRIOR FILING DATE: 2000-07-28  
; PRIOR APPLICATION NUMBER: PCT/US00/03062  
; PRIOR FILING DATE: 2000-02-08  
; PRIOR APPLICATION NUMBER: 60/119,468  
; PRIOR FILING DATE: 1999-02-10  
; NUMBER OF SEQ ID NOS: 173  
; SOFTWARE: Patent In Ver. 2.0  
; SEQ ID NO 50  
; TYPE: DNA  
; ORGANISM: Homo sapiens

## US-11-240-769-50

Query Match 10.4%; Score 190.4; DB 9; Length 1094;  
Best Local Similarity 52.2%; Pred. No. 3.7e-34;  
Matches 494; Conservative 0; Mismatches 446; Indels 6; Gaps 3;  
QY 806 CAGGTGTCATGGAGATGTAATTTAGACATCAAGTCAACCCACTCATATTTTCTCTCTGATT 865  
DB 9 CCGGTGTCACGGCGAGCTTTCAGGGCCCCCAGTAGCCCATCATCTCAGCTCCCTGCTG 68  
QY 866 GGTCTTGATGTCAGATATTTGCTGTCTCTCATCGTTTATTTATTTGTTGCAATGATAGAA 925  
DB 69 GACTCAGGCAATTCAGCTGTTCTGATGATCTCTCATCGTCATCTTTGTAGSCATGCTTGG 128  
QY 926 GATTTATATATGAGA---GGGGAATCAATGCTCAGTACAGCCATATTTGTCATGCTGCT 982  
DB 129 ATGCTGTGGCCCTCCAGCGGGGAGCTCTCATGACACACAGCCGTGCTCTCTCTCATGTTTC 188  
QY 983 ACGTCTCCAGTGAATGTTTATTTGAGGAAAGTCTGTATGCTAGACAAGGAGGAAGAGA 1042  
DB 189 ATGGGGGTGTTTGGCGGATTTTCTGTCGCCGCTGTACCGCACTTTAAAGGCCATCCG 248  
QY 1043 TGGATAAAGCAGATGTTTATTTGGGGCATTTCTTATCCAGCTATGTTGTGTGCACTGCC 1102  
DB 249 TGGAGAAGAGAGCCTTCTGTACGGCAACTCTGTACCTGTTGTGTTTTTGGCATCTGC 308  
QY 1103 TTCTTTCATCAATTTCAATAGCCATTTATACATGCTTCAAGAGCCATTTCTTTTGGAAACA 1162  
DB 309 TTCTGATTTGAATTTGCTTCAATTTGGGGAAGCACTCATCAGAGCGGTGCCCTTCCACC 368  
QY 1163 ATGTTGGCCGTTTGTGTCATCTGTTTTTGTATTTCTTCTTAAATCTTCTGTGTGATCA 1222  
DB 369 ATGTTGGCTTGTCTGTGATGTTG---GTTGGGATCTCTCCCTGCCCCCTGCTACTTGGGCTA 427  
QY 1223 ATACTTGGCCGAATCTGTGAGGTCAAGCCCAACTTTTCTTGTGTGTCATGTCATGTCCT 1282  
DB 428 CTACTTGGCTTCCGAAAGCAG---CCATATGACAACCTCTGTGCGCAACCAAGATTCCTC 485  
QY 1283 GCTCTTACCGGAGAAAAATGTTTCAATGAGGCTCGGGTTATTTGTTGCTTGGTGA 1342  
DB 486 CGGCAGATCCCGAGCAGCGGTGTACATGAACCGATTTTGTGGGATCTCTCATGCTGGG 545  
QY 1343 ATTTTACCTTTTGTGTCATCTTTATGAAATGTAATTTTCACTTTCAGCTTTTCTGGGCA 1402  
DB 546 ATCTTGGCCCTTGGCGCCATGTTTCATGAGCTCTTCTTCACTTTCAGTCTATCTGGGAG 605  
QY 1403 TATAAGATCTATTTATGTCATGCTTTCATGATGCTGTGCTGTTTATCTGTGCAATGTC 1462  
DB 606 AATCAGTTCTATTTACCTCTTTGGCTTCTGTTCTTCTTCTTCTCATCTCTGTTATCC 665  
QY 1463 ACTGTCGTGTGACTATTTGTGTGACATATTTTCTTACTAAATGAGAGAGATTACGGTGG 1522  
DB 666 TGTTCACAAATCAGCATCTGTCATGTTTCTTCCAGCTGTGTGAGAGGATTTACCGCTGG 725  
QY 1523 CAATGGCAAGTTTTTCTCTCTGTCATCACTGCAATCTATGTTTACATGATTTCTCTTT 1582  
DB 726 TGGTGGAGAAATTTTCTAGTCTCCGGGGCTCTGCAATTTCTAGCTCTGTTTATGCCATC 785  
QY 1583 TACTACTATTTTTTCAAACAAAGATGATGCTTATTTTCAAACATCAATTTTACTTTGA 1642  
DB 786 TTTTATTTGTTTAAACAGCTGGACATCGTGGAGTTTCACTCCCTCTCTCTCTTCTTGGC 845  
QY 1643 TATATGGCGGTATTTAGCAGCCCTTGGGATTAATGTTGTGGAGGATTTGTTTACATGGA 1702  
DB 846 TACACGGCCCTCATGGTCTTGTCTTCTTGGCTGTAAAGGGTACCATCGGCTTCTATGCA 905  
QY 1703 ACAAGTGCCTTTGTCGGAATAATCTATATGTAATGAAATTTGACT 1748  
DB 906 GCCTACATGTTTGTTCGCAAGATCTATGCTGCTGTGAAGATAGACT 951

## RESULT 8

US-11-240-769-49



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; Sequence 49, Application US/11240769
; Publication No. US20060036089A1
; GENERAL INFORMATION:
; APPLICANT: Soppet et al.
; TITLE OF INVENTION: 33 Human Secreted Proteins
; FILE REFERENCE: P2037P1C2
; CURRENT APPLICATION NUMBER: US/11/240,769
; CURRENT FILING DATE: 2005-10-03
; PRIOR APPLICATION NUMBER: 09/997,131
; PRIOR FILING DATE: 2001-11-30
; PRIOR APPLICATION NUMBER: 09/628,508
; PRIOR FILING DATE: 2000-07-28
; PRIOR APPLICATION NUMBER: PCT/US00/03062
; PRIOR FILING DATE: 2000-02-08
; PRIOR APPLICATION NUMBER: 60/119,468
; PRIOR FILING DATE: 1999-02-10
; NUMBER OF SEQ ID NOS: 173
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 49
; LENGTH: 1821
; TYPE: DNA
; ORGANISM: Homo sapiens
; US-11-240-769-49

Query Match          9.4%; Score 172; DB 9; Length 1821;
Best Local Similarity 52.7%; Pred. No. 8.5e-30; Indels 20; Gaps 6;
Matches 519; Conservative 0; Mismatches 445;

QY 620 CCGTCCTCTTTTCAACATCGGATTCATTTGGTTTCAATTTTCAACTCTCTTCATGATGGTG 679
DB 537 CTGACCATGAGTGACGTCACGATCCACTGGTTTCTATCACTCACTCGTTGTGTGTC 596

QY 680 ATCTCTTGGTGGCTTAGTTTCAATGATTTTATGAGAACATTAAGAAAGATTATGCT 739
DB 597 TTCTTCTGTGAGTATCTCTGAGCATGATATATCAATCCGACCCCTCCGGAAGGACATGGC 656

QY 740 CGGTACAGTAAAGAGGAAGAAATCGATGATATGATAGAGACCTAGGAGATGAATATGGA 799
DB 657 AACTACACAGAGGAGTACATTTGA-----AGACACCATGGAGGAGTCTGGG 704

QY 800 TGGAAACAGGTGCTAGGATGATTTTA-GACCATCAAGTCACCCACTGATATTTTCTC 858
DB 705 TGGAAAGTTGGTCACGCGGAGCTCTCAGGCCCCCCCAGTACCCCATGATCCTCAGCTC 764

QY 859 TCTGATTTGGTCTGGATGTCAGATATTTGCTGTCTCTCATGCTTATTTATTTGTCAT 918
DB 765 CCGTCTGGCTCAGGCAATTCAGCTGTTCTGTATGATCCTCATGCTCATCTTTGTAGCCAT 824

QY 919 GATAGAGATTTATATATCTGAGA---GGGGATCAATGCTCAGTACAGCCATATTTGTCTA 975
DB 825 GCTTGGGATGCTGTGCGCTCCAGCGGGGAGCTCTCATGACACAGCCCTGCTCTCTT 884

QY 976 TGTGCTACGTCCTCAGTGAATGTTATTTTGGAGGAAGTCTGATCTAGACAAAGGAGG 1035
DB 885 CATGTTTCATGGGGGTGTTTGGCGGATTTCTGCTGGCGGTCTGTACCGCACTTTAAAGG 944

QY 1036 AAGAGATGATTAAGCAGATGTTTATTTGGGCAATTCCTTATCCAGCTATGTTGTGG 1095
DB 945 CCATCGGTGGAAGAAAGAGGAGCTTCTGTACGGCAACTCTGTACCCCTGGTGTGTTTTGG 1004

QY 1096 CACTGCTTCTTCATCAATTTATAGCCATTTATTTACATGCTTCAAGAGCCATTCCTTT 1155
DB 1005 CATCTGCTTCTGATTTGATTTGCTTTCATTTGGGGAAGCACTCATCAGAGGGGTGCCCTT 1064

QY 1156 TGGAAACAAATGGTGGCCGTTTGTGTCATCTGTTTTTTTGTATTTCTCTCTAAATCTGT 1215
DB 1065 TCCACCACTGTTGCTCTGCTGTGTCATGTG-GTTTCGGGATCTCCCTGCCCTCTGCTACT 1123

QY 1216 TGGTACAAATCTTGGCGGAATCTGTGAGGTACGCCCACTTCTCTGCTGTGTCATGTC 1275
DB 1124 TGGGCTACTACTTCCGGCTTCGGAAGCAG--CCATATGACAACCCCTGTGGCGCACCA 1181

QY 1276 TGTGCTCTGCTCTATACCGGAAAAAATGTTTCAATGGAGCCTGCGGTATTTGTTTGCCT 1335
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DB 1182 GATTCCTCCGCGAGATCCCGAGCGGGTGGTACATGACCCGATTTGTGGGATCTCTCAT 1241
QY 1336 GGCTGGAAATTTTACCTTTTGGTTCAATCTTATTTGAAATGATTTTCACTTTCAGCTTTT 1395
DB 1242 GGCTGGGATCTTGCC-TTGGCGCCATGTTTCATCGAGCTCTTCTTCACTTTCAGTGTCTAT 1300
QY 1396 CTGGGCAATATAGATCTATTTATGTTCTATGCTTATGCTGCTCATGATGCTGCTGTTATCCTGTG 1455
DB 1301 CTGGGGAATTCAGTTCTTATTTACCTCTTTGGCTTCTCTGTTCTTCTTCACTATCTCTGTT 1360
QY 1456 CATTTGTGACTGCTGTGTGACTATTGTGTGCACATATTTTCTACTAAATGCAAGAGATTA 1515
DB 1361 GGTATCTCTGTTCAAAATCAGCATCTCATGGTGTACTTCCAGCTGTGTGCGAGAGATTA 1420
QY 1516 CCGGTGGCAATGGAAGATTTTCTCTCTGCTGATCAATCTGCAATCTATGTTTACATGTA 1575
DB 1421 CCGCTGGTGTGGAGAAATTTCTTAGTCTCCGGGGGCTCTGCAATCTTACGTCCTGCTGTTTA 1480
QY 1576 TTCCCTTTTACTACTATTTTTCAA 1599
DB 1481 TGGCATCTTTTATTTTCGTTAAACAA 1504

RESULT 9
US-11-240-769-21
; Sequence 21, Application US/11240769
; Publication No. US20060036089A1
; GENERAL INFORMATION:
; APPLICANT: Soppet et al.
; TITLE OF INVENTION: 33 Human Secreted Proteins
; FILE REFERENCE: P2037P1C2
; CURRENT APPLICATION NUMBER: US/11/240,769
; CURRENT FILING DATE: 2005-10-03
; PRIOR APPLICATION NUMBER: 09/997,131
; PRIOR FILING DATE: 2001-11-30
; PRIOR APPLICATION NUMBER: 09/628,508
; PRIOR FILING DATE: 2000-07-28
; PRIOR APPLICATION NUMBER: PCT/US00/03062
; PRIOR FILING DATE: 2000-02-08
; PRIOR APPLICATION NUMBER: 60/119,468
; PRIOR FILING DATE: 1999-02-10
; NUMBER OF SEQ ID NOS: 173
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 21
; LENGTH: 1816
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (504)
; OTHER INFORMATION: n equals a,t,g, or c
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (1405)
; OTHER INFORMATION: n equals a,t,g, or c
; US-11-240-769-21

Query Match          9.3%; Score 169.4; DB 9; Length 1816;
Best Local Similarity 52.0%; Pred. No. 3.4e-29;
Matches 514; Conservative 7; Mismatches 448; Indels 20; Gaps 6;

QY 616 TGAATCGTCTTTTTCACATCGGATTCATTTGGTTTTCATTTTCAATTTTCACTCTCTTCATGAT 675
DB 520 TTACTTCGACCATGAGTGACGTCAGATCCACTGGTTTCTTATCACTTAACTCCGTTGTTGT 579
QY 676 GGTGATCTTCTTGGTGGGCTTAGTTTCAATGATTTTAAATGAGAAATTAAGAAAGATTA 735
DB 580 GGTCTTCTTCTGTCAGGTATCTCTGAGCATGATTAATCATTCGGACCTCCGGAAGGACAT 639
QY 736 TGTCTGGTACAGTAAAGAGGAAGAAATGATGATGATGATGATGATGATGATGATGATGATGATGAT 795
DB 640 TGGCAACTACACAGAGGAGTACATTTGA-----AGACACCATGAGGAGTGC 687
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QY 796 TGAATGAAACAGTGTGATGAGATGATTTA-GACCATCAAGTCACCCACTGATATTTT 854  
Db 688 TGGGTGGAAGTGTGTGACGCGACGCTCTTCAGGCCCCCCCCAGTAGACCCCATCTCTCA 747  
QY 855 CCTCTCTGATGTTGTTCTGGAATGTCAGATATTTGCTGTGCTCTCATCTGTTATTTATTTG 914  
Db 748 GCTCCCTGCTGGCTCAGGATTCAGCTGTTCTGTATGATCCTCACTGCTATTTGTAG 807  
QY 915 CAATGATAGAAATTTATATACAGAGA---GGGATCAATGCTCAGTACAGCATATTTG 971  
Db 808 CCATGCTTGGATGCTGTGCGCTCCAGCGGGAGCTCTCATGACCAACAGCGCTGCTTC 867  
QY 972 TCTATGCTGCTGCTCTCCAGTGAATGTTATTTGGAGGAAGTCTGTATGCTAGACAAG 1031  
Db 868 TCTTCAATGTTATGCGGGGTTTGGCGGATTTCTGTGCGCGCTGTATACCGCACTTTAA 927  
QY 1032 GAGGAAGGAGATGATTAAGACAGATGTTATTTGGGCAATTCCTTATCCAGCTATGTTG 1091  
Db 928 AAGGCCATCGTGGAGAAAGAGGCTTCTGTACGGCACTCTGTACCTGTGTGTT 987  
QY 1092 GTGGCACTGCTTCTCATCAATTCATAGCCATTTATTAACATGTTCAAGAGCCATTC 1151  
Db 988 TTGGCATCTGCTTCTGATTAATGCTTCTATTTGGGGAAGCACTCATCAGGAGCGTGC 1047  
QY 1152 CTTTGGACAATAGTGGCGCTTGTGTCATCTGTTTGTATTTCTTCTCTAAATC 1211  
Db 1048 CTTTCCCAACATGTTGCTGCTGTGCAATGTG-GTTGGGATCTCCCTGCGCCCTCGTC 1106  
QY 1212 TTGCTGTGATCAATGTTGGCGGAAATCTGTGAGTCAAGCCCACTTCTTGTGCTGTCA 1271  
Db 1107 TACTTGGCTACTTCTGCTTCCGGAAGCAG--CCATATGACAACCTTGTGCGGACCA 1164  
QY 1272 ATGCTGTGCTGCTCTATACCGGAGAAAAATGGTTCAATGAGGCTCGCGTTATTTGTT 1331  
Db 1165 ACCAGATTCCTCCGCGAGATCCCGGAGCGGTGTACATGAACCGATTTGTGGGCACTC 1224  
QY 1332 GCTCGGTGGAATTTACCTTTTGGTTCATCTTTATGAAATGTTATTCATCTTCACTG 1391  
Db 1225 TCATGCTGGATCTTGTGCTTCCGCGCCATGTTCACTGAGCTCTTCTTCACTTCACTG 1284  
QY 1392 CTTTCTGGGCATATAAGATCTATTATGCTCATGCTTCAATGCTGCTGCTGTTATCC 1451  
Db 1285 CTATCTGGGAGATCAGTCTATTAATCTCTTGGCTTCTGCTTCTGTTTCACTATCC 1344  
QY 1452 TGTGCAATGTGACTGTGTGCACTATTTGTGTCACATATTTTCTATCAATATGCAAG 1511  
Db 1345 TGGTGGKATCTCTKTCACAAATCAGCATCGTCAATGCTGATGCTGCTGCTGCTGCTG 1404  
QY 1512 -ATTACGGTGGCAATGGAACAGTTTCTCTGCTGTCATCACTGCAATCTATGTTTAC 1570  
Db 1405 NATTACCGTGTGGTGGGAGAAATTCCTAGTCTCCGCGGGCTCTGCAATTCWACGCTG 1464  
QY 1571 ATGATTTCTTTTACTACTATTTTTCAA 1599  
Db 1465 GTTATGCCATCTTTWATTCGTTAACAA 1493

## RESULT 10

US-09-925-065A-724812/c  
; Sequence 724812, Application US/09925065A  
; Publication No. US20040181048A1  
; GENERAL INFORMATION:  
; APPLICANT: Wang, David G.  
; TITLE OF INVENTION: Identification and Mapping of Single  
; Nucleotide Polymorphisms in the Human Genome  
; FILE REFERENCE: 108927.135  
; CURRENT APPLICATION NUMBER: US/09/925, 065A  
; CURRENT FILING DATE: 2001-08-08  
; PRIOR APPLICATION NUMBER: US 60/243,096  
; PRIOR FILING DATE: 2000-10-24  
; PRIOR APPLICATION NUMBER: US 60/252,147  
; PRIOR FILING DATE: 2000-11-20

; PRIOR APPLICATION NUMBER: US 60/250,092  
; PRIOR FILING DATE: 2000-11-30  
; PRIOR APPLICATION NUMBER: US 60/261,766  
; PRIOR FILING DATE: 2001-01-16  
; PRIOR APPLICATION NUMBER: US 60/289,846  
; PRIOR FILING DATE: 2001-05-09  
; NUMBER OF SEQ ID NOS: 957086  
; SOFTWARE: FastSeq for Windows Version 4.0  
; SEQ ID NO 724812  
; LENGTH: 1251  
; TYPE: DNA  
; ORGANISM: Homo sapiens  
; US-09-925-065A-724812

Query Match 7.6%; Score 138; DB 6; Length 1251;  
Best Local Similarity 86.0%; Pred. No. 5.5e-22;  
Matches 153; Conservative 0; Mismatches 25; Indels 0; Gaps 0;  
QY 991 ACTGAATGCTTATTTGGAGGAAGTCTGTATCTAGACAAGGAGGATGGATAAA 1050  
Db 750 ACTAATAATATATATAACTTTTCATAGTTCTTTTCAGGAGGAGATGGATAAA 691  
QY 1051 GCAGATGTTTATTTGGGGCAATTCCTTATCCAGCTATGTTGTCGCACTGCTTCTTCAT 1110  
Db 690 GCAGATGTTTATTTGGGGCAATTCCTTATCCAGCTATGTTGTCGCACTGCTTCTTCAT 631  
QY 1111 CAATTCATAGCAATTTATACCATGCTTCAAGAGCAATTCCTTTGGAACAATGGTG 1168  
Db 630 CAATTCATAGCAATTTATACCATGCTTCAAGAGCAATTCCTTTTGGAAACAATGGTG 573

## RESULT 11

US-10-932-182A-478  
; Sequence 478, Application US/10932182A  
; Publication No. US20060046253A1  
; GENERAL INFORMATION:  
; APPLICANT: NAKAO, YOSHIHIRO  
; APPLICANT: NAKAMURA, NORIHIRO  
; APPLICANT: KODAMA, YUKIO  
; APPLICANT: FUJIMURA, TOMOKO  
; APPLICANT: ASHIKARI, TOSHIHIRO  
; TITLE OF INVENTION: METHODS FOR ANALYZING GENES OF INDUSTRIAL YEASTS  
; FILE REFERENCE: 030685-043  
; CURRENT APPLICATION NUMBER: US/10/932,182A  
; CURRENT FILING DATE: 2004-09-02  
; NUMBER OF SEQ ID NOS: 197023  
; SOFTWARE: PatentIn version 3.3  
; SEQ ID NO 478  
; LENGTH: 2019  
; TYPE: DNA  
; ORGANISM: Saccharomyces pastorianus  
; US-10-932-182A-478

Query Match 6.8%; Score 124.4; DB 7; Length 2019;  
Best Local Similarity 46.4%; Pred. No. 9.5e-19;  
Matches 567; Conservative 0; Mismatches 631; Indels 24; Gaps 4;  
QY 530 GTTCCAAATCTAAATCCAGATGTCATATTCAGTAAATGAAAGAGTCAGATGTGAAA 589  
Db 817 GATAAAGATATGACGTGTATTTTACTACTCCGTCAAAATTCATTGCTTCTGATACAGTT 876  
QY 590 TTTGAAGATCGATTTTGACAAATATCTTGATCGTCTCTTTTCAACATCGGATTCATCG 649  
Db 877 TGGGCTACAGATGGGACAGTATCTACAT-----ATTATGATCCGCAAAATTCATCG 930  
QY 650 TTTTCAAATTTTCACTCTCTCATGATGTTGATCTTCTTGGTGGGCTTACTGTTCAATGAT 709  
Db 931 TTTTCTTTAATTAATTTCTCCATCATCATTTTATCATCATCTGTGTTTATTCTTCT 990  
QY 710 TTAATGAGAACATTAAGAAAGATTTATGCTCGGTACAGTTAAAGAGAGAAATGATGAT 769  
Db 991 ATACTTCGGGCTGTGAGAGTGAATTTTGGCCGCTTATACGAA-----CTTCAC 1038







		Matches 215; Conservative 0; Mismatches 190; Indels 0; Gaps 0;			
Qy	1349	CCTTTTGGTTCAATCTTTATTTGAATGTATTTTCATCTTCACGCTCTTTCTGGGCATATAAG 1408			
Db	1	CMTTTTGGCTGCATCTTTTATACAGCTTTTCTTCATTTCTGAATAGCAATTTGGTCCACCAG 60			
Qy	1409	ATCTATTATGTCTATGGCTTTCATGATGTGTGGTGTCTGGTTATCCTGTGCAATGTGACTGTC 1468			
Db	61	ATGTATTACATGTTTGGTTTCTCTGTTCTGGTGTTTATCAATTTTGGTTATTACCTGTTC 120			
Qy	1469	TGTGTGACTATTGTGTGCACATATTTTCTACTAAATGAGAGAGATTACGGTGGCAATGG 1528			
Db	121	GAAGCAACTATATCTTTTGTCTACTTTCACCTATGTGCAGAGGATTACCAITGGCAGTGG 180			
Qy	1529	ACAAGTTTTCTCTGTGTCATCACTCAACTATATTTTACATGTATTTCTTTTACTAC 1588			
Db	181	CGTTCCTTCTTACCAAGGGCTTTCACAGCTGTTTACTTCTCGATATAGCCATACACTAC 240			
Qy	1589	TATTTTTTCAAAACAAAGATGTATGGCTTATTTCAAAACATCATTTTACTTTTGGATATATG 1648			
Db	241	TTCCTTCTCAAACTGCAGATCAGGGGACAGCAAGTACAACTCTGTACTTTCGGTTTACT 300			
Qy	1649	GGGTATTATTAGCACAGCCTTGGGGATAATGTGTGAGCGGATTTGGTTACATGGGAACAAGT 1708			
Db	301	ATGATAAATGGTTTGTATCTTCTTCTCTTTTACAGGAACAAATGGCTTCTTTGCATGCTTT 360			
Qy	1709	GCCTTTGTCCGAAATCTATACTAATGTGAAATTTGACTAGAGA 1753			
Db	361	TGGTTTGTACCAAAATATACAGTGTGTGAAAGTGGACTGAAGA 405			

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Job time : 590.727 secs



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